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MANUFACTURERS COUNCIL ON COLOR AND APPEARANCE

COLLABORATIVE REFERENCE PROGRAM

COLOR AND APPEARANCE

COLOR AND COLOR DIFFERENCE

REPORT NO. 28



U.S. DEPARTMENT OF COMMERCE

National Bureau of Standards

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NBS COLLABORATIVE REFERENCE PROGRAMS

TAPPI Paper and Board (6 times per year)

Bursting strength	Smoothness
Tearing strength	Surface pick strength
Tensile breaking strength	K & N ink absorption
Elongation to break	Moisture content
Tensile energy absorption	Opacity
Folding endurance	Blue reflectance (brightness)
Stiffness	Specular gloss, 75°
Air resistance	Thickness
Grammage	Concora (flat crush)
	Ring crush

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Mullen burst of linerboard
Concora test of medium

MCCA Color and Appearance (4 times per year)

Gloss at 60°
Color and color difference

CTS Rubber (4 times per year)

Tensile strength, ultimate elongation and tensile stress
Hardness
Mooney viscosity
Vulcanization properties

ASTM Cement (2 times per year)

Chemical (11 chemical components)
Physical (15 characteristics)

AASHTO Bituminous

Asphalt cement (2 times per year)
Cutbacks (once a year)

NBS Collaborative Reference Programs
A05 Technology Building
National Bureau of Standards
Washington, DC 20234

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MANUFACTURERS COUNCIL ON
COLOR AND APPEARANCE

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**COLLABORATIVE REFERENCE PROGRAM
FOR
COLOR AND APPEARANCE**

COLOR AND COLOR DIFFERENCE

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**U. S. DEPARTMENT OF COMMERCE
National Bureau of Standards**

NBSIR 79-1812

INTRODUCTION

This Collaborative Reference Program is sponsored by the Manufacturers Council on Color and Appearance and the National Bureau of Standards. Four times per year, color chip samples are distributed to each participating laboratory. After the data has been returned to and analyzed by NBS, a report (as illustrated by this report) showing the data from all participants is prepared. A plot of the spectrophotometric curves of the samples was provided by Hemmendinger Color Laboratory, Belvedere, New Jersey. Reflectance values and colorimetric data for 45/0 geometry usually provided by NBS were unavailable for this report.

If there are any questions on the notes, the analyses, or the report in general, contact J. Horlick on 301-921-2946.

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KEY TO TABLES

MEAN	The average of individual test determinations.
GRAND MEAN - (GR. MEAN)	The average of the individual laboratory MEANS, excluding laboratories flagged (see column F) with an X, #, or +.
SD OF MEANS -	The standard deviation of the laboratory MEANS about the GRAND MEAN: an index of the among- laboratory precision.
INST CODE -	Code for instrument type and color space used to report measurements, see Table of Instrument Codes.
F -	Flag, is based on ΔE column with the following meaning: 0 - Included in grand mean analysis. * - Included in grand means but results are between two and three standard deviations. The participants should take this as a warning to reexamine testing procedures. X - Excluded from all calculations because ΔE is beyond (3) standard deviations. + - Excluded from all grand mean calculations because lab is using a nonstandard instrument or test method. # - Excluded from all grand mean calculations because data were not understood, data were extreme, late, or in- sufficient information was supplied concerning instrument and test method used.

Note: In addition to flag (F) based on delta E column, it is also possible to have either an X or an * on individual MEANS as follows:

- X - following a MEAN signifies that the mean is greater than 3 SD of MEANS from the Grand Mean. The values for this laboratory have been omitted in the calculations involving the MEAN for the column.
- * - following any of the MEANS signifies that the quantity is greater than 2 but less than 3 of the appropriate standard deviations from the corresponding average. The participant should take this as a warning to reexamine his testing procedures.

ΔE - Total color difference between two samples. In X, Y, Z analysis it is calculated in MacAdams (FMC II) units. For L, a, b analysis it is calculated in Hunter units.

ΔE Calculation

ΔE is calculated in the Color and Color Difference Collaborative Reference Program by the FMC2* equations (X, Y, Z analysis) as follows:

The yellow-blue chromatic difference is

$$\Delta C_1 = K_1 S(P\Delta P + Q\Delta Q)/bD^2 - K_1 \Delta S/b;$$

the lightness difference is

$$\Delta L = 0.279 K_2 (P\Delta P + Q\Delta Q)/aD;$$

and the red-green chromatic difference is

$$\Delta C_3 = K_1 (Q\Delta P - P\Delta Q)/aD.$$

The quantity, D, is an abbreviation,

$$D = (P^2 + Q^2)^{\frac{1}{2}}.$$

$$K_1 = 0.55669 + 0.049434 Y - 0.82575 \cdot 10^{-3} Y^2 + 0.79172 \cdot 10^{-5} Y^3 - 0.30087 \cdot 10^{-7} Y^4,$$

$$K_2 = 0.17548 + 0.027556 Y - 0.57262 \cdot 10^{-3} Y^2 + 0.63893 \cdot 10^{-5} Y^3 - 0.26731 \cdot 10^{-7} Y^4,$$

$$a^2 = 17.3 \cdot 10^{-6} (P^2 + Q^2) / [1 + 2.73 P^2 Q^2 / (P^4 + Q^4)],$$

$$b^2 = 3.098 \cdot 10^{-4} (S^2 + 0.2015 Y^2)$$

$$P = 0.724 X + 0.382 Y - 0.098 Z,$$

$$Q = -0.48 X + 1.37 Y + 0.1276 Z,$$

$$S = 0.686 Z,$$

$$\Delta E = [(\Delta C_1)^2 + (\Delta L)^2 + (\Delta C_3)^2]^{\frac{1}{2}}$$

*Friele-MacAdam-Chickering metric

Notes on Specific Laboratory Results

- C241
 - Reported extreme a value compared to other labs on samples D05 and D06
- C256A, C256B
 - Data received too late for inclusion in Grand Mean Statistics
- C637, C633
 - Data received too late for inclusion in Grand Mean Statistics
- C320, C538
 - Apparent reporting or measurement problem
- C514, C662
 - Used illuminant D-65
- C549
 - Apparent reporting or measurement problem on mean X samples D07 and D08
- C608
 - Reported extreme X and Y values compared to other labs on samples D07 and D08
- C664
 - Reported extreme Y and Z values on all five samples

Notes on Samples

Several labs reported that the sample identification mark on sample W17 showed through in the test area. Care will be taken to insure that this does not happen in the future.

ANALYSIS C70-1 TABLE 1
COLOR & COLOR DIFFERENCE

INSTRUMENT IDENTIFICATION

INST	CODE	INSTRUMENT	COLOR	DATA
			SPACE	CODE
C70AC		ACS SPECTRA SENSOR	X Y Z	9014
C70BL		B+L 505 SPECTROPHOTOMETER	X Y Z	9014
C70CA		CARY 14	X Y Z	9014
C70CD		COLOR EYE SMALL SPHERE	X Y Z	9014
C70CE		COLOR EYE SMALL SPHERE	XXYZ,4V	9016
C70CF		COLOR EYE SMALL SPHERE	XYZ,BaSd4	9017
C70CG		COLOR EYE SMALL SPHERE	XXYZ,Ba	9018
C70CH		COLOR EYE SMALL SPHERE	XYZ,3V	9011
C70CL		COLOR EYE LARGE SPHERE	XXYZ,4V	9016
C70CM		COLOR EYE LARGE SPHERE	XXYZ,Ba	9018
C70CN		COLOR EYE LARGE SPHERE	XYZ,BaSd4	9017
C70DC		DIAG CBROMASCAN SPECTROPHOTOMETER	X Y Z	9014
C70DH		DIAG MATCH SCAN SPECTROPHOTOMETER	X Y Z	9014
C70DK		DIAG/LSCE AUTOMATE	XYZ,BaSd4	9017
C70DL		DIAG/LSCE AUTOMATE	XYZ,3V,4F	9019
C70DM		DIAG/LSCE AUTOMATE	XXYZ,4V	9016
C70DS		DIAG/SSCE AUTOMATE	XXYZ,Ba	9018
C70DT		DIAG/SSCE AUTOMATE	XYZ,BaSd4	9017
C70GA		GARDNER AUTO AC2/AC3	L a b	9013
C70GB		GARDNER AUTO AC2/AC3	X Y Z	9014
C70GC		GARDNER XL-20/XL-30 SERIES	X Y Z	9014
C70GD		GARDNER XL-20/XL-30 SERIES	L a b	9013
C70GE		GE/DIAG/HARDY SPECTROPHOTOMETER	X Y Z	9014
C70GK		GARDNER XL-70	X Y Z	9014
C70GL		GARDNER XL-70	L a b	9013
C70GM		GARDNER MULTIPURPOSE REFLECTOMETER	X Y Z	9014
C70GP		GARDNER XL-200 SERIES	L a b	9013
C70GX		GARDNER XL-10	L a b	9013
C70GY		GARDNER XL-10	X Y Z	9014
C70HA		HUNTER D25A (DA,D1A,D2A)	L a b	9013
C70BB		HUNTER D25A (DA,D1A,D2A)	X Y Z	9014
C70BF		HUNTER D25AA	L a b	9013
C70BG		HUNTER D25AA	X Y Z	9014
C70BM		HUNTER D25M (DM,D1M,D2M)	L a b	9013
C70BN		HUNTER D25M (DM,D1M,D2M)	X Y Z	9014
C70BP		HUNTER D25P (DP,D1P,D2P)	X Y Z	9014
C70BQ		HUNTER D25P (DP,D1P,D2P)	L a b	9013
C70BR		HUNTER D25A (DA,D1A,D2A)	Rd a b	9012
C70BT		HUNTER D54 SPECTROPHOTOMETER	X Y Z	9014
C70BU		HUNTER D54 SPECTROPHOTOMETER	L a b	9013
C70IB		IBM SPECTROPHOTOMETER	X Y Z	9014
C70KC		KCS-18	XXYZ,4V	9016
C70KD		KCS-18	XXYZ,Ba	9018
C70KS		KCS-18	X Y Z	9014
C70KI		KCS-40	X Y Z	9014
C70LS		LERES TRILAC	X Y Z	9014
C70LT		LERES TRILAC	XYZ,3V	9011
C70MD		MACBETH MS2000 SPECTROPHOTOMETER	X Y Z	9014
C70NE		MACBETH MS2000 SPECTROPHOTOMETER	L a b	9013
C70MG		MACBETH MC1010	L a b	9013
C70MH		MACBETH MC1010	X Y Z	9014
C70MS		MARTIN SWEETS	X Y Z	9014
C70NT		MARTIN SWEETS	XXYZ,Ba	9018
C70ND		NEGTEC 220 DU COLOR	R G B	9015
C70NE		NEGTEC 220 DU COLOR	X Y Z	9014
C70SA		SPECIAL INSTRUMENT - INCLUDED	X Y Z	9014
C70SB		SPECIAL INSTRUMENT - INCLUDED	Rd a b	9012
C70SC		SPECIAL INSTRUMENT - INCLUDED	L a b	9013
C70SL		SPECIAL INSTRUMENT - INCLUDED	R G B	9015
C70SP		SPECIAL INSTRUMENT - EXCLUDED	X Y Z	9014
C70SQ		SPECIAL INSTRUMENT - EXCLUDED	Rd a b	9012
C70SR		SPECIAL INSTRUMENT - EXCLUDED	L a b	9013
C70SS		SPECIAL INSTRUMENT - EXCLUDED	R G B	9015
C70ZD		ZEISS DMC25	X Y Z	9014
C70ZE		ZEISS ELREPHG	X Y Z	9014
C70ZF		ZEISS ELREPHG	R G B	9015
C70XX		GIVE INSTRUMENT MAKE+MODEL.	NET SPECIFIED	9020

FORMAT OF COLORIMETRIC (INPUT) DATA

DATA	CODE	COLOR SCALE

9011	X,Y,Z	3 FUNCTION VITROLITE CORRECTION
9012	Rd,a,b	
9013	L,a,b	HUNTER
9014	X,Y,Z	
9015	R,G,B	
9016	X,X',Y,Z	4 FUNCTION VITROLITE CORRECTION
9017	X,Y,Z,	BaSd4 CORRECTION
9018	X,X',Y,Z	BaSd4 CORRECTION
9019	X,Y,Z	4 FUNCTION VITROLITE CORRECTION
9020	(NGN-STD. INST.)	SCALE SPECIFIED WITH DATA)

MCCA COLLABORATIVE REFERENCE PROGRAM
 X, Y, Z SPACE ANALYSIS, NORMAL DATA
 COLOR + COLOR DIFFERENCE

1978-1979

LAB CODE	F	SAMPLE D05			SAMPLE D06			DIFFERENCE D06 - D05			INST CODE	LAB
		MEAN X	MEAN Y	MEAN Z	MEAN X	MEAN Y	MEAN Z	ΔX	ΔY	ΔZ	ΔE	
C157	G	41.93	42.28	76.90	41.53	41.64	76.42	-0.40	-0.64	-0.48	1.58	70GE C157
C244	X	41.84	42.65	76.65	41.98	42.80	76.88	+0.14 *	+0.15 X	+0.23 X	0.26X	70ZE C244
C250	X	42.26	42.65	76.76	42.41	42.75	77.00	+0.16 *	+0.10 X	+0.24 X	0.44X	70ZF C250
C251	X	41.72	42.50	76.29	41.82	42.70	76.40	+0.10 *	+0.20 X	+0.11 *	0.60X	70ZE C251
C253	X	42.38	43.03	77.83	42.53	43.17	76.99	+0.14 *	+0.14 X	+0.84 *	2.11X	70GC C253
C278	X	42.40	43.55	75.75	42.45	43.65	75.75	+0.05	+0.10 X	+0.00 *	0.28X	70NE C278
C396	X	43.75*	40.95X	75.55	43.55*	40.45X	75.10	-0.20	-0.50	-0.45	2.35X	70GY C396
C407	G	42.62	43.46	77.49	42.66	43.25	77.40	+0.04	+0.21	+0.09	1.68	70BL C407
C414	X	41.82	42.84	75.15	41.54	42.34	74.82	-0.28	-0.50	-0.32	1.35X	70MD C414
C416A	G	42.76	43.83	77.90	42.50	43.30	77.40	-0.26	-0.53	-0.51	1.72	70GE C416A
C416B	G	44.12*	45.42X	78.42	43.97X	45.01X	78.20	-0.14	-0.41	-0.22	1.57	70SA C416B
C418	X	41.53	41.47*	75.81	41.49	41.05*	75.55	+0.03	+0.41	+0.26	2.65X	70CE C418
C422	G	41.07	41.57	74.80	41.17	41.79	74.90	+0.10 *	+0.17 *	+0.10 *	1.75	70SA C422
C424	G	42.45	43.24	76.93	42.43	42.98	76.85	+0.01	+0.26	+0.08	1.54*	70CA C424
C428	X	41.66	42.56	77.31	41.61	42.20	77.13	-0.05	-0.37	-0.18	2.04X	70HB C428
C437	G	42.44	42.66	79.98*	42.40	42.36	79.71*	-0.04	-0.30	-0.26	1.78	70CE C437
C444	G	42.44	43.19	77.48	42.24	42.73	77.11	-0.20	-0.47	-0.37	1.73	70GE C444
C451	G	42.18	43.13	76.37	42.23	42.94	76.21	+0.05	+0.20	+0.15	1.72	70AC C451
C453	G	42.64	44.04	77.15	42.63	43.74*	77.02	-0.01	-0.30	+0.13	1.87	70HT C453
C459	G	41.75	42.64	75.82	41.52	42.14	75.41	-0.24	-0.50	-0.42	1.70	70GE C459
C460	G	41.97	42.65	76.73	41.73	42.33	76.44	-0.24	-0.52	-0.30	1.65	70GE C460
C462A	X	41.35	42.45	77.15	41.15	42.05	76.75	-0.20	-0.40	-0.40	1.32X	70HB C462A
C463	G	42.53	43.22	77.45	42.39	42.84	77.15	+0.14	+0.38	+0.30	1.51*	70ZD C463
C469	X	42.83	43.34	77.18	42.04	42.78	76.73	-0.79 X	-0.56	-0.45	3.06X	70GE C469
C470	G	42.33	42.94	77.11	42.02	42.36	76.49	-0.31	-0.57	-0.62	1.81	70DH C470
C472	G	42.61	43.28	77.17	42.40	42.81	76.73	-0.21	-0.47	-0.44	1.74	70ZD C472
C473	G	42.26	42.84	76.57	42.03	42.37	76.14	-0.23	-0.47	-0.43	1.63	70DH C473
C474	G	41.63	42.46	76.19	41.42	41.99	75.79	-0.21	-0.47	-0.40	1.73	70GE C474
C476	G	42.11	42.87	77.02	41.95	42.45	76.69	-0.16	-0.43	-0.33	1.71	70SA C476
C479B	G	42.58	43.77	78.41	42.94	43.48	78.14	-0.04	-0.29	-0.27	1.72	70SA C479B
C480	G	42.25	43.15	77.76	42.06	42.70	77.39	-0.19	-0.45	-0.38	1.67	70HB C480
C481	G	41.65	42.55	77.10	41.55	42.15	76.90	-0.10	-0.40	-0.20	1.86	70HG C481
C483	G	41.38	42.15	75.70	41.04	41.60	75.11	-0.33	-0.55	-0.59	1.52*	70ZF C483
C495	G	42.59	43.48	76.80	42.36	43.00	76.44	-0.22	-0.48	-0.42	1.65	70KS C495
C496A	G	42.60	43.42	77.54	42.35	42.90	77.09	-0.26	-0.51	-0.45	1.68	70GE C496A
C499C	G	43.86*	45.32X	79.64*	43.44*	44.62X	79.03*	-0.41	-0.69	-0.61	1.73	70BL C499C
C503	G	42.44	43.38	77.56	42.25	42.92	77.22	-0.19	-0.45	-0.34	1.66	70GE C503
C508	G	42.13	43.20	76.72	41.99	42.78	76.47	-0.14	-0.42	-0.25	1.72	70GE C508
C511	G	42.12	42.96	76.58	42.04	42.60	76.26	-0.09	-0.35	-0.31	1.85	70DH C511
C521A	G	42.70	43.65	77.91	42.41	43.12	77.44	-0.29	-0.58	-0.47	1.79	70CA C521A
C521B	G	41.12	41.71*	76.11	40.67	41.01*	75.49	-0.45	-0.70	-0.62	1.70	70SA C521B
C522	G	41.42	42.38	75.51	41.16	41.83	74.97	-0.26	-0.55	-0.54	1.90*	70SA C522
C524	G	42.97	43.93	78.08	42.89	43.56	77.86	-0.09	-0.37	-0.23	1.77	70GE C524
C526	G	42.72	43.51	77.13	42.55	43.07	76.63	-0.17	-0.43	-0.31	1.67	70KS C526
C532	G	42.24	43.18	77.13	42.02	42.69	76.77	-0.22	-0.49	-0.36	1.71	70GE C532
C534	G	41.86	42.79	75.21	41.62	42.29	74.83	-0.24	-0.49	-0.38	1.59	70MD C534
C540	G	43.03	43.93	77.74	42.72	43.34	77.22	-0.31	-0.59	-0.51	1.76	70GE C540
C543	G	41.67	42.66	77.58	41.45	42.18	77.16	-0.22	-0.48	-0.42	1.70	70HB C543
C545	G	40.62*	41.54*	74.41	40.37*	41.03*	73.98	-0.25	-0.50	-0.43	1.67	70SA C545
C549	#	42.36	43.03	76.96	42.38	42.80	76.81	-0.02	-0.23	-0.16	1.74	70DH C549
C552	G	41.14	42.04	76.67	41.18	41.82	76.59	-0.04	-0.22	-0.09	1.76	70HN C552
C608	X	40.87	39.50X	71.48X	42.66	41.48	75.57	1.79 X	1.97 X	4.09 X	4.76X	70GC C608
C612	G	41.79	42.66	76.42	41.64	42.23	76.10	-0.16	-0.43	-0.32	1.77	70GE C612
C627	#	42.40	43.35	76.50	41.97	42.68	75.91	-0.43	-0.68	-0.59	1.64	70SA C627
C630	G	42.49	43.30	77.35	42.29	42.84	76.97	-0.19	-0.47	-0.38	1.74	70KS C630
C631A	G	42.38	43.15	76.53	42.10	42.62	76.05	-0.28	-0.53	-0.48	1.62	70AC C631A
C631B	G	42.59	43.48	76.80	42.25	42.87	76.24	-0.34	-0.61	-0.56	1.74	70AC C631B
C632	G	42.17	43.00	76.23	41.78	42.36	75.53	-0.39	-0.64	-0.70	1.74	70AC C632
C634	X	40.45*	40.70X	73.68*	40.26*	40.23X	73.38*	-0.19	-0.47	-0.49	2.03X	70CE C634
C638	G	41.05	42.10	75.25	40.80	41.60	74.80	-0.25	-0.50	-0.45	1.64	70GC C638
C639	G	41.85	42.32	75.98	41.52	41.73	75.41	-0.33	-0.59	-0.57	1.77	70DH C639
C644	G	42.05	43.19	75.32	41.68	42.51	74.69	-0.41	-0.68	-0.63	1.75	70MD C644
C645	G	42.36	43.21	76.54	41.98	42.56	75.96	-0.37	-0.65	-0.58	1.77	70AC C645
C656	G	42.20	43.19	76.42	41.94	42.67	75.97	-0.26	-0.52	-0.44	1.67	70SA C656
C657	G	40.70	43.23	70.17X	40.41*	42.67	69.72X	-0.30	-0.57	-0.44	1.65	70SA C657

LAB C6DE	F	SAMPLE D05			SAMPLE D06			DIFFERENCE D06 - D05			INST C6DE	LAB
		MEAN X	MEAN Y	MEAN Z	MEAN X	MEAN Y	MEAN Z	ΔX	ΔY	ΔZ	ΔE	
C660	6	41.01	43.56	70.64X	40.73	43.00	70.23X	-0.27	-0.56	-0.41	1.72	70AC C660
C662	6	40.38*	42.53	70.15X	40.19*	42.08	69.91X	-0.18	-0.45	-0.24	1.59	70SP C662
C664	#	39.60X	38.01X	21.35X	39.40X	37.62X	21.19X	-0.21	-0.39	-0.16	1.86	70HT C664
C671A	6	41.54	42.45	75.26	41.64	42.30	75.20	+0.10 *	-0.15 *	-0.06	1.83	70DH C671A
C671B	6	41.37	41.95	75.02	41.29	41.63	74.62	-0.08	-0.32	-0.40	1.81	70DH C671B
C671C	6	40.98	41.98	75.16	40.92	41.66	74.99	-0.07	-0.32	-0.18	1.64	70GC C671C
C671D	6	42.19	43.17	78.10	41.90	42.61	77.57	-0.28	-0.55	-0.52	1.75	70HB C671D
C672	6	41.90	42.85	76.00	41.70	42.40	75.60	-0.20	-0.45	-0.40	1.64	70GC C672
C675	6	42.43	43.29	76.69	42.29	42.90	76.35	-0.14	-0.40	-0.34	1.69	70AC C675
C683	6	39.62X	40.57X	74.19*	39.52X	40.22X	73.84*	-0.10	-0.35	-0.35	1.81	70GC C683
C691	6	41.62	42.85	74.73	41.48	42.48	74.38	-0.14	-0.38	-0.36	1.62	70SA C691
C699	6	41.91	42.55	75.30	41.99	42.79	75.28	+0.09 *	-0.16 *	-0.02 *	1.71	70BL C699
C700	6	41.77	42.57	75.71	41.72	42.26	75.49	-0.05	-0.31	-0.23	1.71	70DH C700
GRAND MEANS												
		42.07	42.94	76.08	41.84	42.49	76.31	-0.19	-0.45	-0.37	1.71	
SD OF MEANS												
		.71	.58	1.18	.67	.58	1.18	.13	.13	.17	.08	
INCLUDED LINES FOR THIS MEAN												
		61	59	59	60	59	59	62	62	62	62	

LAB CODE	F	SAMPLE D07			SAMPLE D08			DIFFERENCE D08 - D07			INST CODE	LAB
		MEAN X	MEAN Y	MEAN Z	MEAN X	MEAN Y	MEAN Z	ΔX	ΔY	ΔZ	ΔE	
C157	G	7.57*	11.40	12.00	7.50*	11.47	11.44	-0.07	0.07	-0.56	2.52	70GE C157
C244	G	8.31	11.95	11.96	8.13	11.90	11.16	-0.19	-0.05	-0.80 *	3.02	70ZE C244
C250	G	8.45	12.00	13.11	8.23	11.95	12.28	-0.22 *	-0.05	-0.83 *	3.00	70ZF C250
C251	G	8.34	11.80	12.40	8.22	11.80	11.69	-0.12	0.00	-0.71	2.76	70ZE C251
C253	X	8.30	11.84	11.90	8.15	11.92	11.15	-0.15	0.09	-0.75	3.53X	70GC C253
C278	G	8.80	12.00	13.75X	8.65	12.00	13.10X	-0.15	0.00	-0.65	2.42	70NE C278
C396	G	8.95*	11.610*	11.85	8.90*	11.15*	11.20	-0.05	0.05	-0.65	2.90	70GY C396
C407	G	8.86	12.36	13.00	8.79*	12.41	12.40	-0.07	0.05	-0.60	2.47	70BL C407
C414	G	8.07	11.64	11.97	7.93	11.64	11.29	-0.14	0.00	-0.68	2.71	70ND C414
C416A	G	8.26	11.99	12.25	8.10	11.95	11.50	-0.16	-0.03	-0.75	2.83	70GE C416A
C416B	G	8.36	12.07	11.79	8.23	12.06	11.10	-0.13	-0.01	-0.69	2.75	70SA C416B
C418	G	8.31	11.90	12.46	8.17	11.88	11.76	-0.14	-0.02	-0.70	2.62	70CE C418
C422	G	8.02	11.47	11.46	7.96	11.52	10.83	-0.06	0.05	-0.63	2.83	70SA C422
C424	G	8.57	12.21	12.97	8.51	12.29	12.27	-0.06	0.08	-0.70	2.99	70CA C424
C428	G	8.02	11.64	11.80	7.92	11.65	11.09	-0.11	0.01	-0.71	2.89	70HB C428
C437	G	8.57	12.08	14.03X	8.45	12.08	13.29X	-0.13	0.00	-0.73	2.59	70CE C437
C444	G	8.27	11.86	12.54	8.17	11.89	11.88	-0.10	0.02	-0.66	2.64	70GE C444
C451	G	8.29	11.84	12.30	8.25	11.91	11.73	-0.04	0.07	-0.58	2.57	70AC C451
C453	G	8.66	12.36	12.31	8.83*	12.64*	12.02	-0.17 X	0.28 X	-0.29 X	2.77	70HT C453
C459	G	8.12	11.66	12.18	8.01	11.66	11.49	-0.11	-0.00	-0.69	2.70	70GE C459
C460	G	8.01	11.59	12.31	7.92	11.61	11.66	-0.09	0.02	-0.66	2.66	70GE C460
C462A	G	8.00	11.60	11.60	7.90	11.60	10.90	-0.10	0.00	-0.70	2.88	70HB C462A
C463	G	8.26	11.87	12.53	8.24	11.98	11.94	-0.02	0.11	-0.59	2.79	70ZD C463
C469	G	8.56	12.14	12.89	8.47	12.16	12.26	-0.08	0.02	-0.63	2.48	70GE C469
C470	G	8.14	11.67	12.53	8.09	11.75	11.93	-0.04	0.08	-0.59	2.65	70DH C470
C472	G	8.18	11.81	12.71	8.11	11.86	12.07	-0.07	0.04	-0.64	2.64	70ZD C472
C473	G	8.14	11.65	12.51	8.11	11.73	11.94	-0.03	0.09	-0.57	2.58	70DH C473
C474	G	8.01	11.56	12.03	7.80	11.47	11.24	-0.21 *	-0.09 *	-0.79	2.79	70GE C474
C476	G	7.99	11.58	12.56	7.80	11.52	11.80	-0.19 *	-0.06	-0.76	2.70	70SA C476
C479B	G	8.54	12.18	12.88	8.44	12.19	12.18	-0.10	0.01	-0.71	2.74	70SA C479B
C480	G	8.21	11.87	11.94	8.19	11.96	11.33	-0.02	0.09	-0.61	2.88	70HB C480
C481	G	7.95	11.45	11.60	7.90	11.50	10.90	-0.05	0.05	-0.70	3.18*	70HG C481
C483	G	8.45	11.85	12.40	8.33	11.90	11.69	-0.12	0.05	-0.71	2.97	70ZF C483
C495	G	8.40	12.01	12.88	8.30	12.03	12.20	-0.10	0.02	-0.68	2.66	70KS C495
C496A	G	8.29	11.91	12.44	8.17	11.91	11.75	-0.12	0.00	-0.69	2.69	70GE C496A
C499C	G	8.95*	12.93X	12.55	8.64	12.72*	11.69	-0.32 X	-0.21 X	-0.87 X	2.84	70BL C499C
C503	G	8.19	11.80	12.48	8.06	11.77	11.75	-0.13	-0.03	-0.73	2.71	70GE C503
C508	G	8.29	11.96	12.12	8.20	11.98	11.50	-0.09	0.02	-0.62	2.54	70GE C508
C511	G	8.23	11.82	12.41	8.11	11.81	11.72	-0.13	-0.01	-0.70	2.66	70DH C511
C521A	G	8.38	12.08	12.30	8.30	12.13	11.67	-0.08	0.05	-0.63	2.67	70CA C521A
C521B	G	7.49*	10.95*	11.63	7.54*	11.20	11.25	0.05 X	0.25 X	-0.38 X	2.71	70SA C521B
C522	G	8.38	11.84	12.80	8.28	11.86	12.00	-0.10	0.02	-0.81 *	3.18*	70SA C522
C524	G	8.74	12.38	13.03	8.65	12.43	12.41	-0.09	0.05	-0.62	2.53	70GE C524
C526	G	8.42	11.98	12.95	8.30	12.00	12.25	-0.12	0.02	-0.70	2.70	70KS C526
C532	G	8.03	11.66	12.15	7.93	11.70	11.45	-0.11	0.04	-0.70	2.92	70GE C532
C534	G	8.13	11.56	12.17	7.98	11.54	11.48	-0.14	-0.02	-0.69	2.63	70ND C534
C540	G	8.98*	12.61*	13.13	8.94*	12.70*	12.54*	-0.04	0.08	-0.59	2.55	70GE C540
C543	G	8.10	11.75	11.87	7.96	11.73	11.12	-0.14	-0.02	-0.75	2.89	70BB C543
C545	G	7.76	11.24	11.68	7.60	11.18	10.97	-0.16	-0.06	-0.71	2.67	70SA C545
C549	#	84.05X	11.96	12.67	82.65X	11.94	11.96	-1.40 X	-0.02	-0.71	2.28*	70DH C549
C552	G	7.90	11.42	11.41	7.76	11.40	10.68	-0.14	-0.03	-0.73	2.90	70HN C552
C608	X	12.35X	8.63X	12.22	12.31X	8.48X	11.44	-0.04	-0.15 X	-0.78	6.05X	70GC C608
C612	G	7.77	11.33	11.77	7.64	11.32	11.08	-0.13	-0.01	-0.69	2.75	70GE C612
C627	*	8.40	12.05	12.56	8.22	11.99	11.77	-0.18	-0.06	-0.78	2.81	70SA C627
C630	G	8.38	12.00	12.57	8.28	12.02	11.93	-0.10	0.03	-0.64	2.56	70KS C630
C631A	G	8.35	11.85	12.51	8.23	11.85	11.85	-0.12	-0.00	-0.66	2.54	70AC C631A
C631B	G	8.48	12.04	12.57	8.39	12.07	11.92	-0.10	0.03	-0.65	2.62	70AC C631B
C632	G	8.32	11.83	12.42	8.23	11.86	11.77	-0.09	0.03	-0.65	2.65	70AC C632
C634	G	8.28	11.84	12.27	8.17	11.88	11.56	-0.11	0.04	-0.70	2.92	70CE C634
C638	G	7.90	11.50	11.50	7.80	11.60	10.90	-0.10	0.10	-0.60	3.00	70GC C638
C639	G	8.15	11.60	12.69	8.05	11.62	12.04	-0.10	0.02	-0.65	2.55	70DH C639
C644	G	8.12	11.76	11.79	8.00	11.76	11.11	-0.11	0.00	-0.68	2.74	70MD C644
C645	G	8.43	11.96	12.52	8.23	11.89	11.74	-0.20 *	-0.07	-0.78	2.80	70AC C645
C656	G	8.26	11.88	12.21	8.13	11.87	11.49	-0.13	-0.01	-0.72	2.77	70SA C656
C657	G	8.14	12.02	11.64	8.03	12.04	11.00	-0.11	0.02	-0.64	2.70	70SA C657

LAB CODE	F	SAMPLE D07			SAMPLE D08			DIFFERENCE D08 - D07			INST CODE	LAB
		MEAN X	MEAN Y	MEAN Z	MEAN X	MEAN Y	MEAN Z	ΔX	ΔY	ΔZ		
C660	G	8.28	12.21	11.86	8.21	12.26	11.28	-0.07	0.04	-0.58	2.51	70AC C660
C662	G	7.98	11.79	11.51	7.89	11.83	10.91	-0.09	0.05	-0.60	2.64	70SP C662
C664	#	7.90	9.98X	3.91X	7.87	9.99X	3.73X	-0.03	0.01	-0.18 X	1.32X	70HT C664
C671A	G	8.16	11.72	12.12	8.09	11.77	11.50	-0.07	0.05	-0.62	2.66	70DH C671A
C671B	G	8.07	11.49	12.52	7.99	11.54	11.90	-0.08	0.05	-0.62	2.59	70DH C671B
C671C	G	7.90	11.34	11.48	7.85	11.44	10.86	-0.05	0.10	-0.62	2.95	70GC C671C
C671D	G	8.22	11.86	11.85	8.09	11.88	11.12	-0.13	0.01	-0.73	3.00	70HB C671D
C672	G	8.75	12.20	12.40	8.50	12.10	11.50	-0.25 X	-0.10 *	-0.99 X	3.21*	70GC C672
C675	G	8.37	11.91	12.39	8.25	11.90	11.70	-0.13	-0.00	-0.69	2.66	70AC C675
C683	M	7.62*	11.05*	11.17*	7.54	11.09*	10.59*	-0.08	0.04	-0.58	2.58	70GC C683
C691	G	8.56	11.88	12.61	8.47	11.87	12.01	-0.09	-0.00	-0.60	2.30*	70SA C691
C699	G	8.80	12.48*	12.87	8.68	12.48	12.17	-0.13	0.00	-0.70	2.64	70BL C699
C700	G	8.18	11.71	12.24	8.11	11.79	11.64	-0.06	0.08	-0.60	2.68	70DH C700
GRAND MEANS												
		8.27	11.81	12.26	8.16	11.85	11.59	-0.10	0.02	-0.67	2.73	
SD OF MEANS												
		0.31	0.32	0.47	0.31	0.34	0.46	0.04	0.05	0.06	0.18	
INCLUDED LABS FOR THIS MEAN												
		73	72	71	73	73	71	69	70	69	73	

NCCA COLLABORATIVE REFERENCE PROGRAM
L, a, b SPACE ANALYSIS, NORMAL DATA
COLOR + COLOR DIFFERENCE

1978-1979

LAH CGDE	F	SAMPLE D05			SAMPLE D06			DIFFERENCE D06 - D05			INST CGDE	LAH
		MEAN I	MEAN A	MEAN B	MEAN L	MEAN A	MEAN H	ΔL	ΔA	ΔH	ΔE	
C105	G	65.10	-0.00	-24.30	64.80	.70	-24.55	-0.30	.70	-0.25	.80	70HM C105
C121	G	64.70	-0.10	-25.10	64.30	.55	-25.50*	-0.40	.65	-0.40	.86	70HM C121
C122	G	65.20	-0.40	-24.40	64.70	.40	-24.75	-0.50	.80	-0.35	1.01	70SC C122
C148	G	65.55	-0.25	-24.20	65.20	.40	-24.50	-0.35	.65	-0.30	.80	70SC C148
C150	G	65.79	-0.80	-24.24	65.17	.03	-24.67	-0.63	.84	-0.48 *	1.15	70HA C150
C152	G	65.90	-0.20	-23.90	65.10	.60	-24.60	-0.80 *	.80	-0.70 X	1.33	70HA C152
C166	G	65.40	-0.40	-24.30	65.05	.30	-24.50	-0.35	.70	-0.20	.81	70HA C166
C183	G	65.35	.10	-23.80	65.05	.80	-24.00	-0.30	.70	-0.20	.79	70HA C183
C213	G	64.30*	-0.20	-24.20	64.10*	-0.00	-24.40	-0.20	.20	-0.20	.35	70SC C213
C223	G	65.97	-0.42	-23.61	65.72	.15	-23.80	-0.26	.57	-0.19	.66	70SC C223
C230	G	65.37	.05	-24.27	64.87	.80	-24.69	-0.50	.71	-0.42	.96	70HA C230
C241	#	65.05	-7.25X	-24.30	64.85	-6.55X	-24.50	-0.20	.70	-0.20	.75	70HA C241
C256A	#	64.70	.68	-24.52	64.63	.70	-24.56	-0.07	.03 *	-0.04	.08*	70HM C256A
C256B	#	65.15	.63	-23.04	65.21	.65	-23.04*	-0.06 *	.02 *	-0.00 *	.07*	70HU C256B
C259	G	65.15	-0.70	-24.65	65.10	-0.70*	-24.65	-0.05	.00 *	.00 *	.05*	70SC C259
C262	G	65.33	.53	-23.86	65.36	.49	-23.89	.03 *	-0.05 *	-0.04 *	.07*	70SB C262
C285	G	64.50	.70	-24.70	64.85	.90	-24.90	-0.05	.20	-0.20	.29	70HA C285
C288	G	65.20	.70	-24.80	65.30	.70	-24.70	.10 *	-0.00 *	.10 X	.14*	70HA C288
C291	G	65.20	.60	-24.35	65.05	.60	-24.40	-0.15	.00 *	-0.05	.16	70HA C291
C301	G	64.65	.50	-25.00	64.60	.50	-25.00	-0.05	-0.00 *	-0.00 *	.05*	70HM C301
C317	G	64.60	.90	-24.30	64.60	.90	-24.30	.00	.00 *	.00 *	.00*	70HM C317
C320	#	65.60	.54	23.54X	65.65	.52	23.56X	.05 *	-0.02 *	.02 *	.06*	70HA C320
C325	G	65.61	.87	-23.13	65.35	1.60*	-23.35	-0.27	.73	-0.22	.80	70SB C325
C340	G	65.69	-0.44	-23.04*	65.41	.15	-23.38	-0.29	.59	-0.34	.74	70SC C340
C352	G	65.65	-0.20	-24.35	65.35	.40	-24.60	-0.30	.60	-0.25	.72	70HA C352
C356	G	65.00	-0.30	-24.40	64.70	.40	-24.70	-0.30	.70	-0.30	.82	70HM C356
C380	G	65.50	-0.70	-24.35	65.15	-0.00	-24.60	-0.35	.70	-0.25	.82	70HA C380
C382	G	65.45	.70	-23.80	65.05	1.35	-24.10	-0.40	.65	-0.30	.82	70HA C382
C402	G	65.36	-0.20	-24.15	65.06	.52	-24.42	-0.30	.72	-0.28	.82	70HA C402
C427	G	65.50	-0.00	-24.10	65.25	.75	-24.40	-0.25	.75	-0.30	.85	70HA C427
C440	G	65.76	-0.03	-24.15	65.42	.69	-24.47	-0.34	.71	-0.31	.85	70HA C440
C442	G	65.10	-0.20	-24.20	64.60	.50	-24.60	-0.50	.70	-0.40	.95	70HM C442
C454	G	66.03	1.04	-23.85	65.83*	1.79*	-24.10	-0.20	.75	-0.25	.81	70SC C454
C456	G	65.47	-0.34	-25.03	65.25	.36	-25.26	-0.21	.70	-0.23	.77	70HA C456
C458	G	65.93	-0.37	-24.34	64.68	.26	-24.62	-0.35	.63	-0.29	.78	70HM C458
C475	G	65.58	-0.19	-24.04	65.17	.32	-24.33	-0.41	.51	-0.29	.72	70HA C475
C477	G	65.91	-0.16	-22.97*	65.55	.57	-23.05*	-0.36	.73	-0.29	.86	70HA C477
C494	G	65.95	-0.20	-23.80	65.50	.70	-24.20	-0.45	.90	-0.40	1.08	70HA C494
C496B	G	65.26	-0.05	-24.60	64.85	.67	-24.88	-0.41	.72	-0.28	.87	70GP C496H
C499A	G	65.20	-0.50	-24.25	64.85	.20	-24.60	-0.35	.70	-0.35	.86	70HA C499A
C506	G	65.10	-0.50	-24.60	64.75	.20	-24.80	-0.35	.70	-0.20	.81	70HA C506
C514A	*	65.18	-0.68	-23.07	64.86	.05	-23.34	-0.32	.73	-0.27	.84	70SR C514A
C517	G	65.50	.80	-24.20	65.00	1.60*	-24.50	-0.50	.80	-0.30	.99	70HQ C517
C538	#	66.25*	4.05X	-23.60	65.90*	4.60X	-23.80	-0.35	.55	-0.20	.68	70GX C538
C541	G	65.20	-0.10	-24.20	64.75	.65	-24.55	-0.45	.75	-0.35	.94	70GP C541
C547	G	63.83X	.61	-24.90	63.47X	1.29	-25.15	-0.36	.68	-0.25	.81	70HQ C547
C548	G	65.15	-1.60*	-22.95*	64.88	-0.82*	-23.22*	-0.27	.78	-0.27	.86	70SB C548
C574	G	65.80	.45	-24.05	65.60	1.20	-24.25	-0.20	.75	-0.20	.80	70HQ C574
C576	G	64.88	-0.76	-24.22	64.51	-0.06	-24.56	-0.37	.71	-0.34	.87	70SB C576
C585	G	65.41	-0.37	-24.61	65.03	.37	-24.96	-0.38	.74	-0.35	.90	70SC C585
C600	G	65.55	-1.40*	-22.95*	65.10	-0.70*	-23.20*	-0.45	.70	-0.25	.87	70GD C600
C619	G	65.00	.20	-23.90	64.40	1.00	-24.20	-0.60	.80	-0.30	1.04	70SC C619
C620	G	63.91X	-0.47	-22.03X	63.40X	.31	-22.33X	-0.51	.78	-0.31	.99	70NG C620
C633	#	65.39	.05	-23.88	64.62	.79	-24.45	-0.77 *	.75	-0.57 *	1.22	70HA C633
C648	G	66.30*	.47	-23.19	65.82*	1.12	-23.57	-0.48	.65	-0.38	.89	70HA C648
C674	G	65.55	.10	-24.00	65.25	.80	-24.30	-0.30	.70	-0.30	.82	70HF C674
C677	G	67.01X	-0.63	-24.40	66.81X	.02	-24.57	-0.20	.65	-0.17	.70	70SC C677

GRAND MEANS

65.36

.41

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INCLUDED LABS FOR THIS MEAN

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LAB CODE	F	SAMPLE D07			SAMPLE D08			DIFFERENCE D08 - D07			INST CODE	LAB
		MEAN L	MEAN A	MEAN B	MEAN L	MEAN A	MEAN B	ΔL	ΔA	ΔB	ΔE	
C105	δ	34.20	-17.20	3.40	34.20	-17.80	4.65	.00	-.60	1.25	1.39	70HM C105
C121	δ	33.70	-18.30	3.70	33.65	-19.00	5.00	-.05	-.70	1.30	1.48	70HM C121
C122	δ	34.00	-17.40	3.25	33.90	-18.05	4.50	-.10	-.65	1.25	1.41	70SC C122
C148	δ	34.40	-18.05	3.85	34.35	-18.70	5.10	-.05	-.65	1.25	1.41	70SC C148
C150	X	34.56	-17.95	3.58	35.57X	-19.98X	4.72	1.01 X	-1.99 X	1.14	2.50X	70HA C150
C152	δ	34.60	-17.40	3.60	34.60	-17.90	4.80	.00	-.50 *	1.20	1.30	70HA C152
C166	δ	34.10	-17.35	3.15	34.10	-18.00	4.40	.00	-.65	1.25	1.41	70HA C166
C183	δ	34.15	-18.20	3.70	34.25	-18.85	4.85	.10	-.65	1.15	1.32	70HA C183
C213	δ	33.40*	-18.45*	3.20	33.40*	-18.60	4.40	.00	-.15 X	1.20	1.21*	70SC C213
C223	δ	35.05*	-17.74	3.39	35.09*	-18.43	4.60	.04	-.68	1.20	1.39	70SC C223
C230	δ	34.35	-17.50	3.63	34.33	-18.15	4.83	-.01	-.65	1.20	1.36	70HA C230
C241	#	34.10	-17.90	3.60	34.15	-18.25	4.80	.05	-.35 X	1.20	1.25	70HA C241
C256A	#	33.83	-17.71	3.49	33.82	-18.28	4.71	-.02	-.57	1.22	1.34	70EN C256A
C256B	#	34.34	-17.12	2.61	34.33	-17.76	3.78	-.01	-.64	1.17	1.33	70HU C256B
C259	δ	34.35	-17.15	3.20	34.35	-17.75	4.45	.00	-.60	1.25	1.39	70SC C259
C262	δ	34.65	-17.98	3.27	34.63	-18.73	4.51	-.02	-.75	1.25	1.46	70SH C262
C285	δ	34.20	-17.75	3.00	34.15	-18.45	4.30	-.05	-.70	1.30	1.48	70HA C285
C288	δ	34.40	-17.90	3.50	34.30	-18.60	4.80	-.10	-.70	1.30	1.48	70HA C288
C291	δ	34.45	-17.50	3.75	34.30	-18.20	5.00	-.15	-.70	1.25	1.44	70HA C291
C301	δ	34.00	-17.20	3.30	33.90	-17.90	4.60	-.10	-.70	1.30	1.48	70HM C301
C317	δ	33.90	-17.70	3.60	33.80	-18.40	4.85	-.10	-.70	1.25	1.44	70HM C317
C320	#	34.54	17.00X	3.42	34.95	17.75X	4.62	.01	.75 X	1.20	1.42	70HA C320
C325	δ	34.57	-17.90	3.41	34.57	-18.64	4.56	.00	-.74	1.15	1.37	70SH C325
C340	δ	34.80	-17.28	2.79	34.75	-17.64	3.96	-.05	-.36 X	1.17	1.22*	70SC C340
C352	δ	34.45	-17.85	3.65	34.45	-18.55	4.90	.00	-.70	1.25	1.43	70HA C352
C356	δ	33.90	-17.40	3.30	33.90	-18.10	4.60	.00	-.70	1.30	1.48	70HM C356
C380	δ	34.30	-17.60	3.70	34.20	-18.20	4.95	-.10	-.60	1.25	1.39	70HA C380
C382	δ	34.30	-17.95	3.40	34.25	-18.65	4.70	-.05	-.70	1.30	1.48	70HA C382
C402	δ	34.30	-17.61	3.72	34.42	-18.25	4.95	.12	-.64	1.23	1.39	70HA C402
C427	δ	34.30	-17.50	3.75	34.30	-18.15	5.00	.00	-.65	1.25	1.41	70HA C427
C440	δ	34.50	-17.63	3.65	34.55	-18.26	4.86	.05	-.64	1.21	1.37	70HA C440
C442	δ	33.90	-17.30	2.80	34.00	-17.90	4.00	.10	-.60	1.20	1.35	70HM C442
C454	δ	34.90	-17.80	3.30	35.05	-18.46	4.55	.15	-.66	1.25	1.42	70SC C454
C456	δ	34.15	-17.74	3.46	34.33	-18.25	4.65	.19 *	-.50 *	1.19	1.31	70HA C456
C458	δ	34.01	-17.62	2.81	34.06	-18.16	4.05	.05	-.54	1.24	1.35	70HM C458
C475	δ	34.64	-17.50	2.39	34.53	-18.14	3.64	-.11	-.64	1.25	1.41	70HA C475
C477	δ	34.75	-17.99	2.92	34.73	-18.66	4.13	-.01	-.67	1.21	1.38	70HA C477
C494	δ	34.60	-18.10	2.50	34.65	-18.95	3.70	.05	-.85 *	1.20	1.47	70HA C494
C496B	δ	34.12	-16.74	3.07	34.06	-17.42	4.34	-.05	-.68	1.27	1.43	70GP C496B
C499A	δ	34.05	-17.75	3.40	34.00	-18.35	4.60	-.05	-.60	1.20	1.34	70HA C499A
C506	δ	34.05	-17.50	3.50	34.15	-18.10	4.70	.10	-.60	1.20	1.35	70HA C506
C514A	*	34.24	-17.55	2.57	34.27	-18.16	3.74	.03	-.61	1.17	1.32	70SR C514A
C517	δ	34.20	-16.10X	2.25*	34.45	-16.60X	3.40*	.25 *	-.50 *	1.15	1.28	70HQ C517
C538	#	35.45X	-14.30X	.35X	35.40*	-15.05X	1.70X	-.05	-.75	1.35 *	1.55	70GX C538
C541	δ	34.15	-17.00	3.00	34.20	-17.60	4.20	.05	-.60	1.20	1.34	70GP C541
C547	δ	32.33X	-18.53*	2.85	32.11X	-19.23*	4.12	-.22 *	-.70	1.27	1.46	70HQ C547
C548	δ	33.99	-18.12	3.69	33.91	-18.77	4.91	-.07	-.65	1.23	1.39	70SB C548
C574	δ	34.60	-17.25	2.20*	34.40	-18.10	3.50*	-.20	-.85 *	1.30	1.57*	70HQ C574
C576	δ	33.83	-17.27	3.41	33.78	-17.89	4.65	-.05	-.62	1.24	1.39	70SB C576
C585	δ	34.38	-17.54	3.41	34.38	-18.14	4.64	.00	-.61	1.23	1.37	70SC C585
C600	δ	34.90	-17.70	3.60	34.80	-18.40	4.80	-.10	-.70	1.20	1.39	70GD C600
C619	δ	34.10	-17.70	2.80	34.00	-18.40	3.90	-.10	-.70	1.10 *	1.31	70SC C619
C620	δ	32.96X	-17.25	2.85	32.80X	-17.93	4.02	-.16	-.69	1.17	1.37	70MG C620
C633	#	34.15	-17.43	3.63	34.22	-18.02	4.83	.07	-.59	1.21	1.34	70HA C633
C648	δ	35.39X	-16.78	4.02	35.35*	-17.60	5.30	-.05	-.82 *	1.28	1.52	70HA C648
C674	δ	34.55	-17.45	3.50	34.40	-18.10	4.75	-.15	-.65	1.25	1.42	70HF C674
C677	δ	34.63	-16.40*	2.27*	34.44	-17.07*	3.54*	-.19	-.67	1.27	1.45	70SC C677
GRAND MEANS												
		34.30	-17.59	3.26	34.31	-18.24	4.49	-.02	-.66	1.23	1.40	
SD OF MEANS												
		.34	.43	.44	.37	.43	.45	.10	.07	.05	.07	
INCLUDED LABS FOR THIS MEAN												
		46	48	49	47	48	49	49	47	49	49	

EXPLANATION OF DATA FOR WHITE SAMPLE

Specimens of a white sample were distributed to the participants along with the usual two pairs of colored specimens, and each participant was asked to return measurement data for the white specimen, reporting results in the same manner as for the colored specimens.

As a first step, three laboratories were selected to serve as "reference" laboratories for the purposes of this analysis and the average of their X, Y, Z values for the white sample were computed. Next, the ratios of the participants data to the combined reference laboratory values were calculated for each participant (transformed to X, Y, Z space if necessary). These ratios are shown in the White Sample Analysis tables.

Two observations can be made about the data in the White Sample Analysis tables. First, the participants as a whole tend to be high compared with the combined average values obtained by the selected reference laboratories. Second, a few participants had noticeably extreme values for one or more of the components and these participants especially should look to the cause.

Next, the ratios in the White Sample Analysis tables were used to "adjust" the data of the normal data tables to obtain the adjusted data table values. The adjustment consisted of dividing the X, Y, Z values of the normal data tables by the respective ratios in the White Sample Analysis tables.

The significant change in the adjusted data tables is in the SD OF MEANS. Comparison of these among-laboratory standard deviations with those in the normal data tables, shows considerable reduction for X, Y, Z. Thus part, at least, of the disagreement among participants is due to errors in standardization that could be corrected through use of an agreed-upon white standard. There is no similar significant change for ΔX , ΔY , and ΔZ .

MCCA COLLABORATIVE REFERENCE PROGRAM
WHITE SAMPLE ANALYSIS
X, Y, Z LABORATORIES

1978-1979

LAB CGDE	RATIO--(LAE/COMBINED)			INST CGDE	PERCENT FROM COMBINED		
	X	Y	Z		X	Y	Z
C157	1.0110	1.0055	1.0137	70GE	1.10	.95	1.37
C244	1.0056	1.0071	1.0141	70ZE	.96	.71	1.41
C250	1.0078	1.0022	1.0055	70ZF	.78	.22	.55
C251	1.0118	1.0077	1.0123	70ZE	1.18	.77	1.23
C253	1.0219	1.0185	1.0205	70GC	2.19	1.85	2.05
C278	1.0368	1.0363	1.0339	70NE	3.68	3.63	3.39
C396	1.0226	.9751	1.0092	70GY	2.26	-2.49	.02
C407	1.0166	1.0147	1.0225	70HL	1.66	1.47	2.25
C414	.9933	.9939	.9876	70MD	.67	-.61	-1.24
C416A	1.0240	1.0228	1.0236	70GE	2.40	2.28	2.36
C416B	1.0577	1.0561	1.0292	70SA	5.77	5.61	2.92
C418	.5915	.9866	1.0071	70CE	-.85	-1.34	.71
C422	.5983	.9533	.9960	70SA	-.17	-.67	-.40
C424	1.0119	1.0114	1.0185	70CA	1.19	1.14	1.85
C428	1.0113	1.0070	1.0169	70HB	1.13	-.70	1.69
C437	1.0174	1.0134	1.0727	70CE	1.74	1.34	7.27
C444	1.0148	1.0152	1.0211	70GE	1.48	1.52	2.11
C451	1.0136	1.0123	1.0061	70AC	1.36	1.33	.61
C453	1.0145	1.0136	1.0094	70HT	1.45	1.36	.94
C459	1.0003	1.0001	1.0010	70GE	.03	.01	.10
C460	1.0102	1.0105	1.0120	70GE	1.02	1.09	1.20
C462A	1.0113	1.0082	1.0233	70HB	1.13	.82	2.33
C463	1.0145	1.0109	1.0209	70ZD	1.45	1.09	2.09
C469	1.0179	1.0167	1.0289	70GE	1.79	1.67	2.89
C470	1.0139	1.0140	1.0222	70DH	1.39	1.40	2.22
C472	1.0035	1.0013	1.0094	70ZD	.35	.13	.94
C473	1.0055	1.0070	1.0095	70DH	.95	.70	.95
C474	1.0073	1.0081	1.0099	70GE	.73	.81	.99
C476	.9391	.9351	.9374	70SA	-.6.09	-6.49	-6.26
C479B	1.0292	1.0291	1.0343	70SA	2.92	2.91	3.43
C480	1.0131	1.0104	1.0185	70HB	1.31	1.04	1.85
C481	1.0102	1.0071	1.0146	70BG	1.02	.71	1.46
C483	1.0022	.9989	1.0038	70ZF	.22	-.11	.38
C495	1.0163	1.0160	1.0156	70KS	1.63	1.60	1.56
C496A	1.0229	1.0224	1.0268	70GE	2.29	2.24	2.68
C499C	1.0311	1.0317	1.0371	70BL	3.11	3.17	3.71
C503	1.0171	1.0168	1.0224	70GE	1.71	1.68	2.24
C508	1.0128	1.0134	1.0152	70GE	1.28	1.34	1.52
C511	1.0061	1.0075	1.0101	70DH	.81	.75	1.01
C521A	1.0155	1.0202	1.0232	70CA	1.95	2.02	2.32
C521B	1.0063	1.0036	1.0138	70SA	.63	.36	1.38
C522	1.0014	1.0008	.9597	70SA	.14	.08	-.03
C524	1.0243	1.0236	1.0276	70GE	2.43	2.36	2.76
C526	1.0175	1.0156	1.0184	70KS	1.75	1.56	1.84
C532	1.0169	1.0154	1.0198	70GE	1.69	1.94	1.98
C534	.9949	.9955	.9872	70MD	-.51	-.45	-1.28
C540	1.0231	1.0226	1.0272	70GE	2.31	2.26	2.72
C543	1.0055	1.0068	1.0185	70BB	.95	.68	1.85
C545	.9705	.9711	.9766	70SA	-.2.95	-2.89	-2.34
C549	1.0108	1.0078	1.012	70DH	1.08	.72	-89.88
C552	.9983	.9957	1.0068	70HN	-.17	-.43	.68
C608	1.0240	.9727	.9960	70GC	2.40	-2.73	-.40
C612	1.0110	1.0122	1.0081	70GE	1.10	1.22	.81
C627	1.0036	1.0021	1.0004	70SA	.36	.21	.04
C630	1.0059	1.0081	1.0150	70KS	.99	.81	1.50
C631A	1.0191	1.0175	1.0141	70AC	1.91	1.75	1.41
C631B	1.0159	1.0149	1.0082	70AC	1.59	1.49	.82
C632	1.0126	1.0110	1.0071	70AC	1.26	1.10	.71
C634	.9576	.9821	.9861	70CE	-.1.24	-1.79	-1.39
C638	1.0006	.9985	.9997	70GC	.06	-.11	-.03
C639	1.0020	.9955	1.0058	70DH	.20	-.05	.58
C644	1.0030	1.0048	.9935	70ND	.30	.48	-.65
C645	1.0158	1.0148	1.0116	70AC	1.58	1.48	1.16
C656	1.0023	1.0011	1.0013	70SA	.23	.11	.13
C657	.9824	1.0126	.9288	70SA	-.1.76	1.26	-7.12
C660	.9572	1.0169	.9324	70AC	-.1.28	1.69	-6.76
C662	.9686	.9915	.9219	70SP	-.3.14	-.85	-7.81
C664	1.0610	.9417	.2865	70HT	6.10	-5.83	-71.35
C671A	.9522	.9914	.9881	70DH	-.78	-.86	-1.19
C671B	.9538	.9920	.9928	70DH	-.62	-.80	-.72
C671C	.9548	.9926	.9902	70GC	-.52	-.74	-.98
C671D	1.0125	1.0101	1.0191	70BB	1.25	1.01	1.91
C672	1.0023	1.0000	.9997	70GC	.23	-.00	-.03
C675	1.0141	1.0126	1.0089	70AC	1.41	1.26	.89
C683	1.0159	1.0162	1.0111	70GC	1.99	1.62	1.11
C691	1.0030	.9964	1.0053	70SA	.30	-.36	.53
C699	.9560	.9979	.9923	70BL	-.40	-.21	-.77
C700	.9577	.9965	.9954	70DH	-.23	-.35	-.46

NCCA COLLABORATIVE REFERENCE PROGRAM
WHITE SAMPLE ANALYSIS
L, a, b LABORATORIES

1978-1979

LAB CGDE	RATIO--(LAB/COMBINED)			INST CGDE	PERCENT FROM COMBINED		
	X	Y	Z		X	Y	Z
C105	1.0170	1.0139	1.0227	70HM	1.070	1.39	2.27
C121	1.0055	1.0013	1.0176	70HM	.55	.13	1.76
C122	1.0100	1.0076	1.0163	70SC	1.00	.76	1.63
C148	1.0167	1.0139	1.0204	70SC	1.67	1.39	2.04
C150	1.0113	1.0056	1.0158	70HA	1.13	.96	1.58
C152	1.0159	1.0129	1.0177	70HA	1.59	1.29	1.77
C166	1.0115	1.0097	1.0184	70HA	1.15	.97	1.84
C183	1.0073	1.0055	1.0079	70HA	.73	.55	.79
C213	.5544	.9908	.9976	70SC	-.56	-.92	-.24
C223	1.0201	1.0156	1.0247	70SC	2.01	1.56	2.47
C230	1.0083	1.0050	1.0141	70HA	.83	.50	1.41
C241	1.0159	1.0023	1.0093	70HA	.59	.23	.93
C256A	1.0022	.9997	1.0057	70HM	.22	-.03	.57
C256B	.5573	.9955	.9945	70HU	-.27	-.41	-.55
C259	1.0149	1.0118	1.0198	70SC	1.49	1.18	1.98
C262	1.0165	1.0138	1.0240	70SB	1.69	1.38	2.40
C285	1.0176	1.0034	1.0151	70HA	.76	.34	1.51
C288	1.0165	1.0150	1.0222	70HA	1.65	1.50	2.22
C291	1.0126	1.0108	1.0164	70HA	1.26	1.08	1.64
C301	1.0016	.9992	1.0108	70HM	.16	-.08	1.08
C317	1.0022	.9992	1.0069	70HM	.22	-.08	.69
C320	1.0231	1.0150	1.0193	70HA	2.31	1.50	1.93
C325	1.0130	1.0110	1.0125	70SB	1.30	1.10	1.25
C340	1.0122	1.0095	1.0158	70SC	1.22	.95	1.58
C352	1.0174	1.0150	1.0246	70HA	1.74	1.50	2.46
C356	1.0046	1.0034	1.0104	70HM	.46	.34	1.04
C380	1.0158	1.0139	1.0726	70HA	1.58	1.39	7.26
C382	1.0117	1.0087	1.0142	70HA	1.17	.87	1.42
C402	1.0081	1.0060	1.0122	70HA	.81	.60	1.22
C427	1.0138	1.0108	1.0164	70HA	1.38	1.08	1.64
C440	1.0135	1.0107	1.0163	70HA	1.39	1.07	1.63
C442	1.0034	1.0013	1.0114	70HM	.34	.13	1.14
C454	1.0161	1.0125	1.0253	70SC	1.61	1.25	2.53
C456	1.0122	1.0054	1.0276	70HA	1.22	.94	2.76
C458	1.0041	1.0014	1.0127	70HM	.41	.14	1.27
C475	1.0116	1.0075	1.0179	70HA	1.16	.75	1.79
C477	1.0195	1.0154	1.0178	70HA	1.95	1.54	1.78
C494	1.0328	1.0288	1.0489	70HA	3.28	2.88	4.89
C496B	1.0244	1.0225	1.0218	70GP	2.44	2.25	2.18
C499A	1.0058	1.0034	1.0151	70HA	.58	.34	1.51
C506	1.0114	1.0087	1.0173	70HA	1.14	.87	1.73
C514A	.5980	.9963	.9959	70SR	-.20	-.37	-.41
C517	1.0368	1.0331	1.0367	70HQ	3.68	3.31	3.67
C538	.5806	.9887	.9839	70GX	-.194	-.113	-.161
C541	1.0285	1.0267	1.0302	70GP	2.85	2.67	3.02
C547	1.0052	1.0028	1.0057	70HQ	.52	.28	.57
C548	.9658	.9658	.9665	70SB	-.3.42	-.3.42	-.3.35
C574	1.0380	1.0331	1.0344	70HQ	3.80	3.31	3.44
C576	1.0013	.9965	1.0077	70SB	.13	-.15	.77
C585	1.0104	1.0075	1.0217	70SC	1.04	.75	2.17
C600	1.0010	.9992	.9984	70GD	.10	-.08	-.16
C619	1.0151	1.0161	1.0171	70SC	1.91	1.61	1.71
C620	.5557	.9611	.9482	70NG	-.4.03	-.3.89	-.5.18
C633	1.0069	1.0047	1.0118	70HA	.69	.47	1.18
C648	1.0184	1.0137	1.0247	70HA	1.84	1.37	2.47
C674	1.0117	1.0087	1.0158	70HF	1.17	.87	1.58
C677	1.0321	1.0265	1.0418	70SC	3.21	2.85	4.18

NCCA COLLABORATIVE REFERENCE PROGRAM
X, Y, Z SPACE ANALYSIS, ADJUSTED DATA
COLOR + COLOR DIFFERENCE

1978-1979

LAB CODE	F	SAMPLE D05			SAMPLE D06			DIFFERENCE D06 - D05			INST CODE LAB		
		MEAN X	MEAN Y	MEAN Z	MEAN X	MEAN Y	MEAN Z	ΔX	ΔY	ΔZ	ΔE	CODE	LAB
C157	G	41.47	41.68*	75.86	41.08	41.25*	75.39	-0.40	-0.63	-0.47	1.58	70GE	C157
C244	X	41.44	42.35	75.59	41.58	42.50	75.81	+0.14 *	+0.15 X	+0.23 X	+0.25X	70ZE	C244
C250	X	41.93	42.56	76.34	42.08	42.66	76.58*	+0.16 *	+0.10 X	+0.23 X	+0.44X	70ZF	C250
C251	X	41.23	42.18	75.36	41.33	42.37	75.47	+0.10 *	+0.20 X	+0.11 *	+0.60X	70ZE	C251
C253	X	41.48	42.25	76.26	41.62	42.39	75.44	+0.14 *	+0.14 X	+0.82 *	+2.05X	70GC	C253
C278	X	40.90*	42.02	73.27X	40.94	42.12	73.27X	+0.05	+0.10 X	+0.00 *	+0.28X	70NE	C278
C396	X	42.78X	41.55	75.53	42.59X	41.48	75.08	-0.20	-0.51	-0.45	+2.27X	70GY	C396
C407	G	41.93	42.63	75.78	41.97	42.62	75.70	+0.04	+0.21	+0.09	+1.67	70BL	C407
C414	X	42.10	43.10	76.09	41.83	42.60	75.77	+0.28	+0.50	+0.33	+1.35X	70MD	C414
C416A	G	41.76	42.65	76.11	41.50	42.33	75.62	+0.26	+0.52	+0.49	+1.71	70GE	C416A
C416B	G	41.71	43.01	76.20	41.57	42.62	75.98	+0.14	+0.39	+0.22	+1.52*	70SA	C416B
C418	X	41.88	42.03	75.28	41.85	41.61	75.02	+0.03	+0.42	+0.25	+2.66X	70CE	C418
C422	G	41.14	42.25	75.10*	41.24	42.08	75.20	+0.10 *	+0.18 *	+0.10 *	+1.75	70SA	C422
C424	G	41.65	42.76	75.54	41.94	42.50	75.46	+0.01	+0.26	+0.08	+1.54	70CA	C424
C426	X	41.20	42.27	76.03	41.15	41.91	75.85	+0.04	+0.36	+0.18	+2.04X	70HB	C428
C437	G	41.71	42.10	74.55X	41.67	41.80	74.31*	+0.04	+0.30	+0.24	+1.80	70CE	C437
C444	G	41.82	42.55	75.89	41.63	42.09	75.52	+0.19	+0.46	+0.37	+1.73	70GE	C444
C451	G	41.62	42.57	75.91	41.66	42.38	75.75	+0.04	+0.19	+0.15	+1.71	70AC	C451
C453	G	42.03	43.46*	76.43	42.02	43.15*	76.30	+0.01	+0.30	+0.13	+1.85	70HT	C453
C459	G	41.75	42.64	75.82	41.52	42.14	75.41	+0.24	+0.50	+0.42	+1.70	70GE	C459
C460	G	41.55	42.39	75.83	41.31	41.88	75.53	+0.24	+0.51	+0.29	+1.64	70GE	C460
C462A	X	40.89*	42.10	75.39	40.69*	41.71	75.00	+0.20	+0.40	+0.39	+1.32X	70HB	C462A
C463	G	41.93	42.76	75.67	41.78	42.38	75.58	+0.14	+0.38	+0.29	+1.51*	702D	C463
C469	X	42.08	42.63	75.01*	41.30	42.08	74.58*	+0.78 X	+0.55	+0.43	+3.04X	70GE	C469
C470	G	41.76	42.35	75.44	41.45	41.78	74.84	+0.31	+0.57	+0.61	+1.80	70DH	C470
C472	G	42.46*	43.23	76.46	42.25*	42.76	76.02	+0.21	+0.47	+0.44	+1.74	702D	C472
C473	G	41.87	42.55	75.85	41.64	42.07	75.42	+0.23	+0.47	+0.43	+1.62	70DH	C473
C474	G	41.33	42.13	75.44	41.12	41.66	75.05	+0.20	+0.47	+0.40	+1.73	70GE	C474
C476	G	44.84X	45.85X	82.17X	44.67X	45.40X	81.82X	+0.17	+0.45	+0.35	+1.74	70SA	C476
C479B	G	41.76	42.53	75.82	41.72	42.25	75.55	+0.04	+0.28	+0.26	+1.71	70SA	C479B
C480	G	41.70	42.71	76.35	41.51	42.27	75.98	+0.19	+0.45	+0.37	+1.66	70HB	C480
C481	G	41.23	42.25	75.99	41.13	41.85	75.79	+0.10	+0.40	+0.29	+1.85	70HG	C481
C483	G	41.25	42.20	75.42	40.95	41.65	74.83	+0.33	+0.55	+0.59	+1.52*	70ZF	C483
C495	G	41.91	42.60	75.68	41.69	42.33	75.27	+0.22	+0.47	+0.41	+1.64	70KS	C495
C496A	G	41.65	42.47	75.52	41.40	41.97	75.08	+0.25	+0.50	+0.44	+1.67	70GE	C496A
C499C	G	42.54*	43.93X	76.79*	42.13	43.26*	76.20	+0.40	+0.67	+0.59	+1.72	70BL	C499C
C503	G	41.73	42.66	75.87	41.54	42.22	75.53	+0.19	+0.45	+0.33	+1.66	70GE	C503
C508	G	41.60	42.63	75.57	41.46	42.21	75.33	+0.14	+0.41	+0.24	+1.71	70GE	C508
C511	G	41.79	42.64	75.82	41.70	42.29	75.50	+0.08	+0.35	+0.31	+1.84	70DH	C511
C521A	G	41.89	42.83	76.15	41.60	42.27	75.69	+0.28	+0.56	+0.46	+1.78	70CA	C521A
C521B	G	40.87*	41.57X	75.08*	40.42X	40.87X	74.46*	+0.45	+0.70	+0.61	+1.70	70SA	C521B
C522	G	41.37	42.35	75.54	41.10	41.80	74.99	+0.26	+0.54	+0.55	+1.89*	70SA	C522
C524	G	41.96	42.92	75.99	41.87	42.56	75.77	+0.08	+0.36	+0.22	+1.76	70GE	C524
C526	G	41.99	42.64	75.74	41.82	42.42	75.45	+0.17	+0.43	+0.30	+1.66	70KS	C526
C532	G	41.54	42.36	75.63	41.33	41.88	75.28	+0.22	+0.49	+0.35	+1.70	70GE	C532
C534	G	42.08	42.96	76.19	41.83	42.48	75.81	+0.25	+0.50	+0.38	+1.59	70MD	C534
C540	G	42.06	42.96	75.68	41.76	42.39	75.18	+0.30	+0.57	+0.50	+1.75	70GE	C540
C543	G	41.28	42.37	76.17	41.06	41.89	75.76	+0.22	+0.48	+0.41	+1.70	70HB	C543
C545	G	41.86	42.78	76.20	41.60	42.26	75.76	+0.26	+0.52	+0.44	+1.69	70SA	C545
C549	#	41.91	42.70	76.03X	41.93	42.47	75.81X	+0.02	+0.23	+1.53 X	+0.27X	70DH	C549
C552	G	41.21	42.22	76.10	41.25	42.01	76.07	+0.05	+0.22	+0.08	+1.76	70HN	C552
C608	X	39.92X	40.61X	71.77X	41.66	42.64	75.87	+0.75 X	+0.03 X	+4.11 X	+4.69X	70GC	C608
C612	G	41.34	42.15	75.81	41.19	41.73	75.49	+0.15	+0.42	+0.32	+1.76	70GE	C612
C627	#	42.25	43.27	76.47	41.83	42.59	75.88	+0.42	+0.67	+0.58	+1.64	70SA	C627
C630	G	42.07	42.96	76.21	41.88	42.50	75.83	+0.19	+0.46	+0.37	+1.73	70KS	C630
C631A	G	41.58	42.41	75.47	41.31	41.89	74.99	+0.27	+0.52	+0.48	+1.60	70AC	C631A
C631B	G	41.93	42.84	76.18	41.59	42.24	75.62	+0.34	+0.60	+0.56	+1.72	70AC	C631B
C632	G	41.65	42.54	75.69	41.26	41.90	75.00	+0.39	+0.64	+0.69	+1.73	70AC	C632
C634	X	40.96*	41.45X	74.92*	40.77*	40.97X	74.42*	+0.19	+0.48	+0.50	+2.03X	70CE	C634
C638	G	41.03*	42.15	75.27	40.78*	41.65	74.82	+0.25	+0.50	+0.45	+1.63	70GC	C638
C639	G	41.77	42.34	75.54	41.44	41.75	74.98	+0.33	+0.59	+0.57	+1.77	70DH	C639
C644	G	41.97	42.95	75.82	41.56	42.31	75.19	+0.41	+0.67	+0.63	+1.75	70MD	C644
C645	G	41.70	42.58	75.67	41.33	41.94	75.09	+0.37	+0.64	+0.58	+1.76	70AC	C645
C656	G	42.10	43.14	76.32	41.84	42.62	75.88	+0.26	+0.52	+0.44	+1.67	70SA	C656
C657	G	41.43	42.70	75.55	41.13	42.14	75.07	+0.30	+0.56	+0.48	+1.64	70SA	C657

MCCA COLLABORATIVE REFERENCE PROGRAM
X, Y, Z SPACE ANALYSIS, ADJUSTED DATA
COLOR + COLOR DIFFERENCE

1978-1979

LAB CODE	F	SAMPLE D05			SAMPLE D06			DIFFERENCE D06 - D05			INST CODE	LAB
		MEAN X	MEAN Y	MEAN Z	MEAN X	MEAN Y	MEAN Z	ΔX	ΔY	ΔZ		
C660	G	41.54	42.84	75.77	41.26	42.29	75.33	-0.28	-0.55	-0.44	1.70	70AC C660
C662	G	41.69	42.90	76.10	41.50	42.45	75.83	-0.19	-0.45	-0.27	1.58	70SP C662
C664	#	37.33X	40.37X	74.53X	37.13X	39.95X	73.95X	-0.19	-0.41	-0.58	1.53	70HT C664
C671A	G	41.87	42.83	76.17	41.97	42.67	76.11	+0.10 *	-0.16 *	-0.06	1.83	70DH C671A
C671B	G	41.63	42.29	75.57	41.55	41.97	75.16	-0.08	-0.32	-0.41	1.81	70DH C671B
C671C	G	41.20	42.29	75.91	41.13	41.97	75.73	-0.07	-0.32	-0.18	1.63	70GC C671C
C671D	G	41.67	42.74	76.64*	41.39	42.19	76.12	-0.28	-0.55	-0.52	1.75	70HB C671D
C672	G	41.80	42.65	76.02	41.60	42.40	75.62	-0.20	-0.45	-0.40	1.64	70GC C672
C675	G	41.85	42.76	76.01	41.71	42.37	75.68	-0.14	-0.39	-0.34	1.68	70AC C675
C683	G	38.85X	39.92X	73.37X	38.75X	39.58X	73.03X	-0.09	-0.34	-0.35	1.79	70GC C683
C691	G	41.49	43.01	74.34X	41.36	42.63	73.99X	-0.13	-0.38	-0.35	1.62	70SA C691
C699	G	42.08	43.04	75.89	42.16	42.88	75.87	+0.09 *	-0.17 *	-0.03 *	1.71	70BL C699
C700	G	41.87	42.73	76.06	41.82	42.41	75.84	-0.06	-0.31	-0.23	1.71	70DH C700
GRAND MEANS												
		41.71	42.64	75.87	41.54	42.21	75.48	-0.19	-0.45	-0.36	1.70	
SD OF MEANS												
		.32	.32	.35	.32	.38	.42	.13	.13	.17	.08	
INCLUDED LABS FOR THIS MEAN												
		60	58	58	59	59	59	62	62	62	62	

NCCA COLLABORATIVE REFERENCE PROGRAM
X, Y, Z SPACE ANALYSIS, ADJUSTED DATA
CGLCR + CGLCR DIFFERENCE

1978-1979

LAB CODE	F	SAMPLE D07			SAMPLE D08			DIFFERENCE D08 - D07			INST CODE	LAB
		MEAN X	MEAN Y	MEAN Z	MEAN X	MEAN Y	MEAN Z	ΔX	ΔY	ΔZ	ΔE	
C157	G	7.49*	11.29	11.84	7.42*	11.36	11.29	-0.07	0.07	-0.55	2.50	70GE C157
C244	G	8.24	11.87	11.79	8.05	11.82	11.01	-0.18	-0.05	-0.79	3.01	70ZE C244
C250	G	8.38	11.97	13.04	8.17	11.92	12.22	-0.21 *	-0.05	-0.82 *	3.00	70ZF C250
C251	G	8.24	11.71	12.25	8.12	11.71	11.55	-0.12	0.00	-0.70	2.75	70ZE C251
C253	X	8.12	11.63	11.66	7.98	11.71	10.93	-0.15	0.08	-0.73	3.50X	70GC C253
C278	G	8.49	11.58	13.30*	8.34	11.58	12.67*	-0.14	0.00	-0.63	2.39	70NE C278
C396	G	8.75	11.38	11.85	8.70*	11.43	11.20	-0.05	0.05	-0.65	2.91	70GY C396
C407	G	8.71	12.19	12.72	8.65	12.23	12.13	-0.06	0.05	-0.59	2.45	70BL C407
C414	G	8.12	11.71	12.12	7.98	11.71	11.44	-0.14	0.00	-0.68	2.72	70ND C414
C416A	G	8.07	11.72	11.97	7.91	11.69	11.23	-0.16	-0.03	-0.74	2.81	70GE C416A
C416B	G	7.91	11.43	11.46	7.79	11.42	10.78	-0.12	-0.01	-0.68	2.71	70SA C416B
C418	G	8.38	12.06	12.37	8.24	12.04	11.68	-0.14	-0.02	-0.69	2.62	70CE C418
C422	G	8.04	11.55	11.51	7.98	11.60	10.88	-0.06	0.05	-0.63	2.83	70SA C422
C424	G	8.47	12.08	12.73	8.41	12.16	12.05	-0.06	0.08	-0.69	2.97	70CA C424
C428	G	7.94	11.56	11.60	7.83	11.57	10.91	-0.10	0.01	-0.69	2.87	70HB C428
C437	G	8.43	11.92	13.08	8.30	11.92	12.39	-0.13	0.00	-0.68	2.55	70CE C437
C444	G	8.15	11.69	12.28	8.06	11.71	11.64	-0.10	0.02	-0.65	2.62	70GE C444
C451	G	8.18	11.68	12.23	8.14	11.75	11.66	-0.04	0.07	-0.57	2.57	70AC C451
C453	G	8.54	12.20	12.20	8.71*	12.48*	11.91	-0.17 X	0.28 X	-0.29 X	2.76	70BT C453
C459	G	8.12	11.66	12.18	8.01	11.66	11.49	-0.11	-0.00	-0.69	2.70	70GE C459
C460	G	7.93	11.47	12.17	7.84	11.49	11.52	-0.08	0.02	-0.65	2.64	70GE C460
C462A	G	7.91	11.51	11.34	7.81	11.51	10.65	-0.10	0.00	-0.68	2.86	70HB C462A
C463	G	8.15	11.75	12.28	8.13	11.85	11.70	-0.02	0.10	-0.58	2.77	70ZD C463
C469	G	8.41	11.94	12.53	8.33	11.96	11.92	-0.08	0.02	-0.61	2.45	70GE C469
C470	G	8.03	11.51	12.26	7.98	11.59	11.68	-0.04	0.08	-0.58	2.62	70DH C470
C472	G	8.16	11.80	12.60	8.09	11.84	11.96	-0.07	0.04	-0.63	2.64	70ZD C472
C473	G	8.07	11.57	12.39	8.03	11.65	11.83	-0.03	0.08	-0.56	2.57	70DH C473
C474	G	7.95	11.47	11.91	7.75	11.38	11.13	-0.20 *	-0.09 *	-0.78	2.78	70GE C474
C476	G	8.51	12.39*	13.40*	8.31	12.32	12.59*	-0.20 *	-0.07	-0.81 *	2.77	70SA C476
C479B	G	8.29	11.83	12.46	8.20	11.84	11.77	-0.10	0.01	-0.68	2.71	70SA C479B
C480	G	8.10	11.75	11.72	8.08	11.84	11.13	-0.02	0.09	-0.59	2.86	70BB C480
C481	G	7.87	11.37	11.43	7.82	11.42	10.74	-0.05	0.05	-0.69	3.16*	70EG C481
C483	G	8.43	11.86	12.35	8.31	11.91	11.65	-0.12	0.05	-0.71	2.97	70ZF C483
C495	G	8.27	11.82	12.68	8.17	11.85	12.02	-0.09	0.02	-0.66	2.64	70KS C495
C496A	G	8.10	11.65	12.12	7.99	11.65	11.45	-0.11	0.00	-0.67	2.67	70GE C496A
C499C	G	8.68	12.53*	12.11	8.38	12.33	11.27	-0.31 X	-0.20 X	-0.83 *	2.81	70BL C499C
C503	G	8.06	11.61	12.21	7.93	11.58	11.50	-0.13	-0.02	-0.71	2.69	70GE C503
C508	G	8.19	11.81	11.94	8.10	11.83	11.33	-0.09	0.02	-0.61	2.52	70GE C508
C511	G	8.17	11.74	12.29	8.04	11.73	11.60	-0.12	-0.01	-0.69	2.65	70DH C511
CS21A	G	8.22	11.84	12.02	8.15	11.89	11.41	-0.07	0.05	-0.62	2.65	70CA CS21A
CS21B	G	7.44*	10.91*	11.47	7.49*	11.16*	11.10	0.05 X	0.25 X	-0.37 X	2.71	70SA CS21B
CS22	G	8.37	11.84	12.81	8.27	11.86	12.00	-0.10	0.02	-0.81 *	3.18*	70SA CS22
CS24	G	8.53	12.10	12.68	8.45	12.15	12.08	-0.08	0.05	-0.60	2.51	70GE CS24
CS26	G	8.28	11.80	12.72	8.16	11.82	12.03	-0.12	0.02	-0.68	2.68	70KS CS26
CS32	G	7.90	11.44	11.92	7.80	11.48	11.23	-0.10	0.04	-0.69	2.90	70GE CS32
CS34	G	8.17	11.62	12.33	8.03	11.59	11.63	-0.15	-0.03	-0.70	2.64	70MD CS34
CS40	G	8.78*	12.34	12.78	8.74*	12.42*	12.21	-0.04	0.08	-0.57	2.53	70GE CS40
CS43	G	8.03	11.68	11.65	7.89	11.65	10.92	-0.14	-0.02	-0.73	2.87	70HB CS43
CS45	G	8.00	11.57	11.97	7.83	11.52	11.23	-0.16	-0.06	-0.73	2.70	70SA CS45
CS49	*	83.15X	11.87	12.50.22X	81.77X	11.85	11.81.15X	-1.39 X	-0.02	-7.06 X	3.38X	70DB CS49
CS52	G	7.92	11.47	11.34	7.78	11.45	10.61	-0.14	-0.03	-0.73	2.90	70HN CS52
C608	X	12.06X	8.28X	12.27	12.02X	8.72X	11.49	-0.04	-0.16 X	-0.78	6.13X	70GC C608
C612	G	7.69	11.20	11.68	7.56	11.19*	10.99	-0.13	-0.01	-0.69	2.74	70GE C612
C627	#	8.37	12.03	12.55	8.20	11.97	11.77	-0.18	-0.06	-0.78	2.80	70SA C627
C630	G	8.30	11.90	12.38	8.20	11.93	11.75	-0.10	0.02	-0.63	2.55	70KS C630
C631A	G	8.20	11.65	12.34	8.08	11.65	11.69	-0.12	-0.00	-0.66	2.53	70AC C631A
C631B	G	8.35	11.86	12.47	8.26	11.89	11.82	-0.09	0.03	-0.64	2.61	70AC C631B
C632	G	8.22	11.70	12.33	8.13	11.73	11.69	-0.09	0.03	-0.65	2.64	70AC C632
C634	G	8.38	12.06	12.44	8.27	12.10	11.73	-0.11	0.04	-0.71	2.94	70CE C634
C638	G	7.90	11.51	11.50	7.80	11.61	10.90	-0.10	0.10	-0.60	3.00	70GC C638
C639	G	8.14	11.61	12.62	8.04	11.63	11.98	-0.10	0.02	-0.65	2.55	70DB C639
C644	G	8.10	11.70	11.87	7.98	11.70	11.19	-0.11	0.00	-0.68	2.74	70MD C644
C645	G	8.30	11.79	12.38	8.11	11.72	11.61	-0.19 *	-0.06	-0.77	2.78	70AC C645
C656	G	8.25	11.87	12.20	8.12	11.86	11.48	-0.13	-0.01	-0.72	2.77	70SA C656
C657	G	8.29	11.88	12.54	8.17	11.90	11.85	-0.12	0.02	-0.69	2.73	70SA C657

LAB CGDE	F	SAMPLE D07			SAMPLE D08			DIFFERENCE D08 - D07			INST CGDE	LAB
		MEAN X	MEAN Y	MEAN Z	MEAN X	MEAN Y	MEAN Z	ΔX	ΔY	ΔZ	ΔE	
C660	G	8.39	12.01	12.72	8.32	12.06	12.10	-0.08	.04	-0.62	2.56	70AC C660
C662	G	8.24	11.89	12.49	8.15	11.94	11.83	-0.09	.05	-0.65	2.70	70SP C662
C664	#	7.45*	10.60X	13.65*	7.42*	10.61X	13.02*	-0.03	.01	-0.63	2.70	70HT C664
C671A	G	8.23	11.82	12.27	8.16	11.88	11.64	-0.07	.06	-0.62	2.67	70DH C671A
C671B	G	8.12	11.59	12.62	8.04	11.64	11.99	-0.08	.05	-0.62	2.60	70DH C671B
C671C	G	7.94	11.42	11.59	7.90	11.53	10.97	-0.05	.10	-0.62	2.96	70GC C671C
C671D	G	8.12	11.75	11.63	7.99	11.76	10.91	-0.13	.01	-0.72	2.95	70HB C671D
C672	G	8.73	12.20	12.40	8.48	12.10	11.50	-0.25 X	-0.10 *	-0.90 X	3.21*	70GC C672
C675	G	8.26	11.76	12.28	8.14	11.76	11.60	-0.12	-0.00	-0.68	2.65	70AC C675
C683	G	7.47*	10.87*	11.05*	7.40*	10.91X	10.47*	-0.07	.04	-0.57	2.57	70GC C683
C691	G	8.54	11.92	12.54	8.44	11.92	11.95	-0.09	-0.01	-0.60	2.29*	70SA C691
C699	G	8.84*	12.51*	12.97	8.71*	12.51*	12.27	-0.13	.00	-0.70	2.65	70BL C699
C700	G	8.20	11.75	12.30	8.13	11.83	11.70	-0.07	.08	-0.60	2.68	70DH C700
GRAND MEANS												
		8.20	11.75	12.22	8.10	11.78	11.56	-0.10	.02	-0.67	2.72	
SD OF MEANS												
		.28	.30	.48	.28	.28	.48	.04	.05	.06	.18	
INCLUDED LABS FOR THIS MEAN												
		73	73	73	73	72	73	69	70	70	73	

LAB CODE	F	SAMPLE D05			SAMPLE D06			DIFFERENCE D06 - D05			INST CODE LAH		
		MEAN L	MEAN A	MEAN B	MEAN L	MEAN A	MEAN B	ΔL	ΔA	ΔB	ΔE	CODE	LAH
C105	#	64.65	-0.34	-23.54	64.35	.36	-23.78	-0.30	.69	-0.25	.80	70HM	C105
C121	0	64.66	-0.57	-23.96	64.26	.08	-24.36	-0.40	.65	-0.40	.86	70HM	C121
C122	0	64.95	-0.67	-23.71	64.46	.13	-24.06	-0.50	.80	-0.35	1.00	70SC	C122
C148	0	65.10	-0.56	-23.59	64.75	.09	-23.89	-0.35	.65	-0.30	.79	70SC	C148
C150	0	65.48	-0.59	-23.65	64.85	-0.15	-24.13	-0.62	.84	-0.48 *	1.15	70HA	C150
C152	0	65.48	-0.54	-23.42	64.68	.25	-24.11	-0.79 *	.80	-0.69 *	1.32	70HA	C152
C166	0	65.09	-0.60	-23.59	64.74	.09	-23.79	-0.35	.70	-0.20	.80	70HA	C166
C183	0	65.17	-0.11	-23.57	64.87	.59	-23.77	-0.30	.70	-0.20	.78	70HA	C183
C213	0	64.60	-0.60	-23.84	64.40	-0.40	-24.04	-0.20	.20	-0.20	.35	70SC	C213
C223	0	65.46	-0.91	-22.82	65.21	-0.34	-23.00	-0.25	.57	-0.19	.65	70SC	C223
C230	0	65.21	-0.28	-23.58	64.71	.43	-24.00	-0.49	.71	-0.42	.96	70HA	C230
C241	#	64.97	-7.62X	-23.79	64.77	-6.93X	-23.99	-0.20	.70	-0.20	.75	70HA	C241
C256A	#	64.71	.40	-24.11	64.64	.42	-24.15	-0.07	.03 *	-0.04	.08*	70HM	C256A
C256H	#	65.28	.48	-23.19	65.35	.50	-23.19	.07 *	.02 *	-0.00 *	.07*	70HU	C256H
C259	0	64.77	-1.03	-23.96	64.72	-1.03*	-23.96	-0.05	.00 *	-0.00 *	.05*	70SC	C259
C262	0	64.89	.18	-23.01	64.92	.13	-23.04	.03 *	-0.05 *	-0.04 *	.07*	70SH	C262
C285	0	64.79	.22	-23.85	64.74	.42	-24.05	-0.05	.20	-0.20	.28	70HA	C285
C288	0	64.72	.52	-24.12	64.82	.52	-24.02	.10 *	-0.00 *	.10 X	.14*	70HA	C288
C291	0	64.85	.39	-23.84	64.70	.39	-23.89	.15	.00 *	-0.05	.16	70HA	C291
C301	0	64.68	.23	-24.21	64.63	.23	-24.21	-0.05	.00 *	-0.00 *	.05*	70HM	C301
C317	0	64.63	.56	-23.78	64.63	.56	-23.78	.00	.00 *	.00 *	.00*	70HM	C317
C320	#	65.11	-0.37	23.46X	65.16	-0.39	23.48X	.05 *	-0.02 *	.02 *	.06*	70HA	C320
C325	0	65.26	.64	-22.90	64.99	1.36*	-23.12	-0.27	.72	-0.22	.80	70SH	C325
C340	0	65.38	-0.74	-22.51	65.10	-0.16	-22.85	.28	.59	-0.34	.73	70SC	C340
C352	0	65.16	-0.47	-23.52	64.87	.12	-23.76	.30	.60	-0.25	.71	70HA	C352
C356	0	64.89	-0.43	-23.87	64.59	.26	-24.17	-0.30	.70	-0.30	.82	70HM	C356
C380	0	65.05	-0.90	-20.37X	64.70	-0.20	-20.62X	-0.35	.70	-0.25	.82	70HA	C380
C382	0	65.17	.35	-23.32	64.77	1.00	-23.62	.40	.65	-0.30	.82	70HA	C382
C402	0	65.16	-0.44	-23.65	64.87	.27	-23.93	-0.30	.71	-0.27	.82	70HA	C402
C427	0	65.15	-0.34	-23.59	64.90	.40	-23.89	-0.25	.74	-0.30	.84	70HA	C427
C440	0	65.41	-0.39	-23.64	65.07	.31	-23.95	-0.34	.71	-0.31	.84	70HA	C440
C442	0	65.06	-0.44	-23.49	64.56	.26	-23.89	-0.50	.70	-0.40	.95	70HM	C442
C454	0	65.62	.62	-22.83	65.42*	1.36*	-23.08	-0.20	.74	-0.25	.80	70SC	C454
C456	0	65.16	-0.66	-23.67	64.95	.04	-23.90	-0.21	.70	-0.23	.76	70HA	C456
C458	0	64.98	-0.67	-23.54	64.63	-0.04	-23.83	-0.34	.63	-0.28	.78	70HM	C458
C475	0	65.34	-0.66	-23.24	64.93	.15	-23.53	-0.41	.51	-0.29	.71	70HA	C475
C477	0	65.40	-0.61	-22.43*	65.05	.11	-22.71*	-0.36	.72	-0.28	.86	70HA	C477
C494	0	65.02	-0.64	-22.14*	64.58	.25	-22.54*	-0.44	.89	-0.39	1.07	70HA	C494
C496B	0	64.54	-0.26	-24.38	64.14	.45	-24.66	-0.41	.71	-0.28	.86	70GP	C496H
C499A	0	65.09	-0.77	-23.41	64.74	-0.07	-23.76	-0.35	.70	-0.35	.86	70HA	C499A
C506	0	64.82	-0.80	-23.90	64.47	-0.11	-24.10	-0.35	.70	-0.20	.80	70HA	C506
C514A	*	65.30	-0.87	-23.14	64.99	-0.15	-23.41	-0.32	.73	-0.27	.84	70SR	C514A
C517	0	64.44	.38	-23.57	63.95*	1.16	-23.86	-0.49	.79	-0.30	.97	70HQ	C517
C538	#	66.63X	5.08X	-24.08	66.27X	5.63X	-24.28	-0.35	.55	-0.20	.69	70GX	C538
C541	0	64.35	-0.30	-23.65	63.90*	.44	-23.99	-0.44	.74	-0.35	.93	70GP	C541
C547	0	63.75X	.34	-24.66*	63.38X	1.02	-24.91*	-0.36	.68	-0.25	.81	70HQ	C547
C548	0	66.30X	-1.63*	-23.30	66.02X	-0.84*	-23.58	-0.27	.79	-0.27	.88	70SH	C548
C574	0	64.74	-0.10	-23.58	64.54	.63	-23.77	-0.20	.74	-0.20	.79	70HQ	C574
C576	0	64.93	-1.08	-23.60	64.56	-0.37	-23.94	-0.38	.71	-0.34	.87	70SH	C576
C585	0	65.16	-0.70	-23.55	64.79	.04	-23.89	-0.37	.74	-0.35	.90	70SC	C585
C600	0	65.58	-1.60*	-23.01	65.13	-0.90*	-23.26	-0.45	.70	-0.25	.87	70GD	C600
C619	0	64.48	-0.14	-23.64	63.89*	.65	-23.94	-0.60	.79	-0.30	1.04	70SC	C619
C620	0	65.19	-0.31	-23.40	64.67	.49	-23.71	-0.53	.80	-0.31	1.01	70MG	C620
C633	#	65.24	-0.21	-23.34	64.47	.54	-23.90	-0.77 *	.74	-0.57 *	1.21	70HA	C633
C648	0	65.85*	-0.06	-22.30*	65.37	.59	-22.67*	-0.48	.65	-0.37	.89	70HA	C648
C674	0	65.27	-0.24	-23.41	64.97	.45	-23.71	-0.30	.70	-0.30	.81	70HF	C674
C677	0	66.07*	-1.02	-23.16	65.87X	-0.38	-23.33	-0.20	.64	-0.17	.69	70SC	C677
GRAND MEANS													
		65.06	-0.38	-23.49	64.71	.22	-23.75	-0.32	.60	-0.26	.75		
SD OF MEANS													
		.37	.53	.50	.33	.50	.48	.17	.26	.11	.30		
INCLUDED LABS FOR THIS MEAN													
		48	50	49	47	50	49	50	50	48	50		

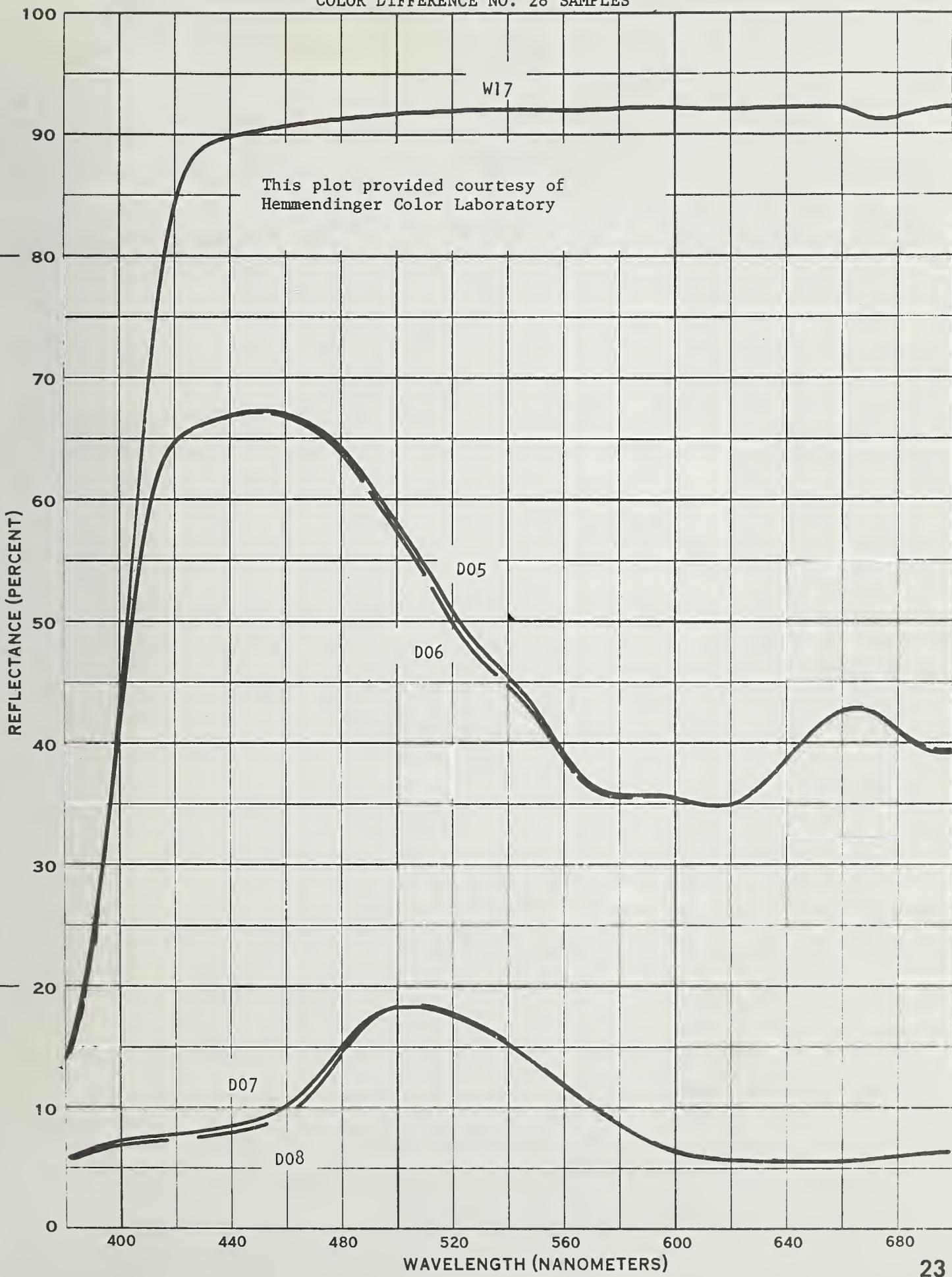
LAB CODE	F	SAMPLE D07			SAMPLE D08			DIFFERENCE D08 - D07			INST CODE	LAB
		MEAN I	MEAN A	MEAN B	MEAN L	MEAN A	MEAN B	AL	AA	AB		
C105	G	33.96	-17.21	3.55	33.96	-17.80	4.78	.00	-.59	1.23	1.37	70HM C105
C121	G	33.68	-18.46	4.02	33.63	-19.15	5.29	-.05	-.70	1.28	1.46	70HM C121
C122	G	33.87	-17.43	3.41	33.77	-18.08	4.65	-.10	-.65	1.23	1.40	70SC C122
C148	G	34.16	-18.04	3.95	34.11	-18.68	5.18	-.05	-.64	1.23	1.39	70SC C148
C150	X	34.40	-17.97	3.69	35.40X	-19.95X	4.82	1.00 X	-1.98 X	1.13	2.49X	70HA C150
C152	G	34.38	-17.42	3.67	34.38	-17.91	4.86	.00	-.50 *	1.19	1.29	70HA C152
C166	G	33.94	-17.34	3.31	33.94	-17.99	4.54	.00	-.65	1.23	1.39	70HA C166
C183	G	34.06	-18.22	3.74	34.16	-18.87	4.88	.10	-.65	1.14	1.32	70HA C183
C213	G	33.55	-18.68*	3.35	33.55	-18.83	4.55	.00	-.15 X	1.20	1.21*	70SC C213
C223	G	34.77*	-17.80	3.55	34.81	-18.47	4.74	.04	-.68	1.19	1.37	70SC C223
C230	G	34.26	-17.60	3.80	34.25	-18.24	4.99	-.01	-.64	1.19	1.35	70HA C230
C241	#	34.06	-18.03	3.74	34.11	-18.38	4.93	.05	-.35 X	1.19	1.24	70HA C241
C256A	#	33.84	-17.82	3.61	33.82	-18.38	4.82	-.02	-.57	1.21	1.34	70HM C256A
C256B	#	34.41	-17.21	2.58	34.40	-17.85	3.75	-.01	-.64	1.17	1.33	70HU C256B
C259	G	34.15	-17.18	3.34	34.15	-17.77	4.58	.00	-.59	1.23	1.37	70SC C259
C262	G	34.41	-17.99	3.45	34.39	-18.73	4.68	-.02	-.74	1.23	1.43	70SB C262
C285	G	34.14	-17.90	3.24	34.09	-18.59	4.52	-.05	-.70	1.28	1.46	70HA C285
C288	G	34.14	-17.83	3.62	34.05	-18.52	4.90	-.10	-.69	1.28	1.46	70HA C288
C291	G	34.27	-17.48	3.84	34.12	-18.18	5.08	-.15	-.69	1.24	1.43	70HA C291
C301	G	34.01	-17.31	3.54	33.91	-18.01	4.82	-.10	-.70	1.28	1.47	70HM C301
C317	G	33.91	-17.63	3.76	33.81	-18.53	5.00	-.10	-.70	1.24	1.43	70HM C317
C320	#	34.68	16.26X	3.48	34.69	17.00X	4.67	.01	.74 X	1.19	1.40	70HA C320
C325	G	34.38	-17.88	3.42	34.38	-18.62	4.57	.00	-.73	1.14	1.36	70SB C325
C340	G	34.63	-17.31	2.91	34.58	-17.67	4.06	-.05	-.35 X	1.15	1.21*	70SC C340
C352	G	34.19	-17.82	3.81	34.19	-18.51	5.04	.00	-.69	1.23	1.41	70HA C352
C356	G	33.84	-17.42	3.44	33.84	-18.12	4.72	.00	-.70	1.29	1.47	70HM C356
C380	G	34.06	-17.55	4.78*	33.96	-18.15	5.95*	-.10	-.59	1.17	1.32	70HA C380
C382	G	34.15	-18.00	3.50	34.10	-18.69	4.79	-.05	-.69	1.29	1.46	70HA C382
C402	G	34.19	-17.65	3.83	34.31	-18.29	5.05	.12	-.64	1.22	1.38	70HA C402
C427	G	34.12	-17.53	3.84	34.12	-18.18	5.08	.00	-.64	1.24	1.39	70HA C427
C440	G	34.32	-17.67	3.75	34.37	-18.30	4.95	.05	-.63	1.20	1.35	70HA C440
C442	G	33.88	-17.38	3.01	33.98	-17.98	4.19	.10	-.60	1.19	1.33	70HM C442
C454	G	34.68	-17.84	3.54	34.83*	-18.50	4.77	.15	-.65	1.23	1.40	70SC C454
C456	G	33.99	-17.78	3.80	34.17	-18.28	4.97	.18 *	-.50 *	1.17	1.28	70HA C456
C458	G	33.99	-17.72	3.04	34.04	-18.26	4.26	.05	-.53	1.22	1.33	70HM C458
C475	G	34.51	-17.62	2.60	34.40	-18.25	3.84	-.10	-.63	1.23	1.39	70HA C475
C477	G	34.48	-18.02	2.95	34.47	-18.68	4.15	-.01	-.66	1.20	1.37	70HA C477
C494	G	34.11	-18.01	2.87	34.16	-18.84	4.04	.05	-.83 *	1.16	1.43	70HA C494
C496B	G	33.74	-16.64*	3.03	33.69	-17.30*	4.28	-.05	-.67	1.25	1.42	70GP C496B
C499A	G	33.99	-17.82	3.63	33.94	-18.42	4.81	-.05	-.60	1.18	1.33	70HA C499A
C506	G	33.90	-17.54	3.66	34.00	-18.13	4.84	.10	-.60	1.19	1.33	70HA C506
C514A	*	34.31	-17.66	2.57	34.33	-18.27	3.74	.03	-.61	1.17	1.32	70SR C514A
C517	G	33.65	-16.00X	2.29*	33.89	-16.49X	3.42*	.25 *	-.49 *	1.13	1.25	70HQ C517
C538	#	35.65X	-13.98X	2.23X	35.60X	-14.74X	1.60X	-.05	-.76	1.36 X	1.56*	70GX C538
C541	G	33.70	-16.86	3.03	33.75	-17.45	4.21	.05	-.59	1.18	1.32	70GP C541
C547	G	32.29X	-18.59*	2.90	32.07X	-19.29*	4.17	-.22 *	-.69	1.26	1.46	70HQ C547
C548	G	34.58	-18.44	3.77	34.51	-19.10	5.01	-.07	-.66	1.25	1.41	70SB C548
C574	G	34.04	-17.18	2.19*	33.85	-18.01	3.47*	-.20	-.83 *	1.28	1.54*	70HQ C574
C576	G	33.86	-17.40	3.60	33.80	-18.02	4.82	-.05	-.62	1.23	1.38	70SB C576
C585	G	34.25	-17.60	3.69	34.25	-18.20	4.90	.00	-.60	1.21	1.35	70SC C585
C600	G	34.91*	-17.78	3.58	34.81	-18.48	4.79	-.10	-.70	1.20	1.39	70GD C600
C619	G	33.83	-17.68	2.80	33.73	-18.38	3.89	-.10	-.69	1.09 *	1.29	70SC C619
C620	G	33.61	-17.52	2.63	33.45	-18.23	3.84	-.16	-.70	1.21	1.41	70MG C620
C633	#	34.07	-17.49	3.76	34.14	-18.07	4.95	.07	-.59	1.19	1.33	70HA C633
C648	G	35.15X	-16.87	4.21	35.10*	-17.68	5.47	-.04	-.81 *	1.26	1.50	70HA C648
C674	G	34.40	-17.50	3.63	34.25	-18.15	4.86	-.15	-.64	1.24	1.40	70HF C674
C677	G	34.15	-16.32X	2.52	33.96	-16.98*	3.75	-.18	-.66	1.24	1.42	70SC C677

GRAND MEANS
 34.12 -17.67 3.42 34.13 -18.28 4.63 -.02 -.66 1.22 1.38

SD OF MEANS
 .31 .42 .50 .35 .46 .51 .10 .07 .04 .07

INCLUDED LABS FOR THIS MEAN
 47 47 49 48 48 49 49 47 49 49

SPECTROPHOTOMETRIC CURVES OF COLOR AND
COLOR DIFFERENCE NO. 28 SAMPLES



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